

Dimitrios Paraskevis

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/558885/publications.pdf>

Version: 2024-02-01

248

papers

8,487

citations

53794

45

h-index

62596

80

g-index

265

all docs

265

docs citations

265

times ranked

9800

citing authors

#	ARTICLE	IF	CITATIONS
1	Full-genome evolutionary analysis of the novel corona virus (2019-nCoV) rejects the hypothesis of emergence as a result of a recent recombination event. <i>Infection, Genetics and Evolution</i> , 2020, 79, 104212.	2.3	544
2	An automated genotyping system for analysis of HIV-1 and other microbial sequences. <i>Bioinformatics</i> , 2005, 21, 3797-3800.	4.1	468
3	Prevalence of Drug-Resistant HIV-1 Variants in Untreated Individuals in Europe: Implications for Clinical Management. <i>Journal of Infectious Diseases</i> , 2005, 192, 958-966.	4.0	385
4	A global scientific strategy to cure hepatitis B. <i>The Lancet Gastroenterology and Hepatology</i> , 2019, 4, 545-558.	8.1	342
5	European guidelines on the clinical management of HIV-1 tropism testing. <i>Lancet Infectious Diseases</i> , The, 2011, 11, 394-407.	9.1	218
6	Transmission of Drug-Resistant HIV-1 Is Stabilizing in Europe. <i>Journal of Infectious Diseases</i> , 2009, 200, 1503-1508.	4.0	213
7	Impact of Hepatitis B Virus Infection on the Progression of AIDS and Mortality in HIV-Infected Individuals: A Cohort Study and Meta-Analysis. <i>Clinical Infectious Diseases</i> , 2009, 48, 1763-1771.	5.8	193
8	The Global Spread of Hepatitis C Virus 1a and 1b: A Phylodynamic and Phylogeographic Analysis. <i>PLoS Medicine</i> , 2009, 6, e1000198.	8.4	177
9	Insights to SARS-CoV-2 life cycle, pathophysiology, and rationalized treatments that target COVID-19 clinical complications. <i>Journal of Biomedical Science</i> , 2021, 28, 9.	7.0	167
10	Transmission of HIV Drug Resistance and the Predicted Effect on Current First-line Regimens in Europe. <i>Clinical Infectious Diseases</i> , 2016, 62, 655-663.	5.8	135
11	Dating the origin and dispersal of hepatitis B virus infection in humans and primates. <i>Hepatology</i> , 2013, 57, 908-916.	7.3	131
12	HIV-1 subtype distribution and its demographic determinants in newly diagnosed patients in Europe suggest highly compartmentalized epidemics. <i>Retrovirology</i> , 2013, 10, 7.	2.0	129
13	Phylogenetic Analysis of SARS-CoV-2 Data Is Difficult. <i>Molecular Biology and Evolution</i> , 2021, 38, 1777-1791.	8.9	119
14	SARS-CoV-2 wastewater surveillance data can predict hospitalizations and ICU admissions. <i>Science of the Total Environment</i> , 2022, 804, 150151.	8.0	116
15	Tracing the HIV-1 subtype B mobility in Europe: a phylogeographic approach. <i>Retrovirology</i> , 2009, 6, 49.	2.0	114
16	Transmission of drug-resistant HIV-1 in Europe remains limited to single classes. <i>Aids</i> , 2008, 22, 625-635.	2.2	109
17	European recommendations for the clinical use of HIV drug resistance testing: 2011 update. <i>AIDS Reviews</i> , 2011, 13, 77-108.	1.0	106
18	Adverse effects of COVID-19 mRNA vaccines: the spike hypothesis. <i>Trends in Molecular Medicine</i> , 2022, 28, 542-554.	6.7	104

#	ARTICLE	IF	CITATIONS
19	Risk of triple-class virological failure in children with HIV: a retrospective cohort study. <i>Lancet</i> , The, 2011, 377, 1580-1587.	13.7	101
20	HIV-1 outbreak among injecting drug users in Greece, 2011: a preliminary report. <i>Eurosurveillance</i> , 2011, 16, .	7.0	100
21	Economic Recession and Emergence of an HIV-1 Outbreak among Drug Injectors in Athens Metropolitan Area: A Longitudinal Study. <i>PLoS ONE</i> , 2013, 8, e78941.	2.5	97
22	The Calculated Genetic Barrier for Antiretroviral Drug Resistance Substitutions Is Largely Similar for Different HIV-1 Subtypes. <i>Journal of Acquired Immune Deficiency Syndromes</i> (1999), 2006, 41, 352-360.	2.1	90
23	Impact of HIV-1 Subtype on CD4 Count at HIV Seroconversion, Rate of Decline, and Viral Load Set Point in European Seroconverter Cohorts. <i>Clinical Infectious Diseases</i> , 2013, 56, 888-897.	5.8	88
24	Analytical methodologies for the detection of SARS-CoV-2 in wastewater: Protocols and future perspectives. <i>TrAC - Trends in Analytical Chemistry</i> , 2021, 134, 116125.	11.4	88
25	Development and assessment of a novel real-time PCR assay for quantitation of HBV DNA. <i>Journal of Virological Methods</i> , 2002, 103, 201-212.	2.1	68
26	Increasing Prevalence of HIV-1 Subtype A in Greece: Estimating Epidemic History and Origin. <i>Journal of Infectious Diseases</i> , 2007, 196, 1167-1176.	4.0	68
27	Molecular epidemiology of HIV-1 infection in Europe: An overview. <i>Infection, Genetics and Evolution</i> , 2016, 46, 180-189.	2.3	68
28	Molecular epidemiology reveals the role of war in the spread of HIV in Ukraine. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 1051-1056.	7.1	65
29	In vivo characteristics of human immunodeficiency virus type 1 intersubtype recombination: determination of hot spots and correlation with sequence similarity. <i>Journal of General Virology</i> , 2003, 84, 2715-2722.	2.9	63
30	Design and baseline findings of a large-scale rapid response to an HIV outbreak in people who inject drugs in Athens, Greece: the ARISTOTLE programme. <i>Addiction</i> , 2015, 110, 1453-1467.	3.3	62
31	Global Dispersal Pattern of HIV Type 1 Subtype CRF01_AE: A Genetic Trace of Human Mobility Related to Heterosexual Sexual Activities Centralized in Southeast Asia. <i>Journal of Infectious Diseases</i> , 2015, 211, 1735-1744.	4.0	62
32	Immunomodulatory Function of HBeAg Related to Short-Sighted Evolution, Transmissibility, and Clinical Manifestation of Hepatitis B Virus. <i>Frontiers in Microbiology</i> , 2018, 9, 2521.	3.5	62
33	Primary resistance to integrase strand-transfer inhibitors in Europe: Table 1.. <i>Journal of Antimicrobial Chemotherapy</i> , 2015, 70, 2885-2888.	3.0	61
34	A network intervention that locates and intervenes with recently HIV-infected persons: The Transmission Reduction Intervention Project (TRIP). <i>Scientific Reports</i> , 2016, 6, 38100.	3.3	60
35	The global spread of HIV-1 subtype B epidemic. <i>Infection, Genetics and Evolution</i> , 2016, 46, 169-179.	2.3	60
36	Molecular characterization of occult hepatitis B cases in Greek blood donors. <i>Journal of Medical Virology</i> , 2009, 81, 815-825.	5.0	59

#	ARTICLE	IF	CITATIONS
37	A review of the impact of weather and climate variables to COVID-19: In the absence of public health measures high temperatures cannot probably mitigate outbreaks. <i>Science of the Total Environment</i> , 2021, 768, 144578.	8.0	59
38	Impact of Antiretroviral Therapy on Tuberculosis Incidence Among HIV-Positive Patients in High-Income Countries. <i>Clinical Infectious Diseases</i> , 2012, 54, 1364-1372.	5.8	58
39	Molecular epidemiology of a hepatitis C virus outbreak in a haemodialysis unit. <i>Nephrology Dialysis Transplantation</i> , 1999, 14, 1188-1194.	0.7	57
40	Integrating Phylodynamics and Epidemiology to Estimate Transmission Diversity in Viral Epidemics. <i>PLoS Computational Biology</i> , 2013, 9, e1002876.	3.2	57
41	Subgenotype A1 of HBV “tracing human migrations in and out of Africa. <i>Antiviral Therapy</i> , 2013, 18, 513-521.	1.0	54
42	Limited cross-border infections in patients newly diagnosed with HIV in Europe. <i>Retrovirology</i> , 2013, 10, 36.	2.0	52
43	Efficient and targeted COVID-19 border testing via reinforcement learning. <i>Nature</i> , 2021, 599, 108-113.	27.8	51
44	Homelessness and Other Risk Factors for HIV Infection in the Current Outbreak Among Injection Drug Users in Athens, Greece. <i>American Journal of Public Health</i> , 2015, 105, 196-204.	2.7	50
45	One year into the pandemic: Short-term evolution of SARS-CoV-2 and emergence of new lineages. <i>Infection, Genetics and Evolution</i> , 2021, 92, 104869.	2.3	49
46	Rapid decline of HIV incidence among people who inject drugs during a fast-track combination prevention programme following an HIV outbreak in Athens. <i>Journal of Infectious Diseases</i> , 2017, 215, 1496-1505.	4.0	48
47	Recent HIV-1 Outbreak Among Intravenous Drug Users in Romania: Evidence for Cocirculation of CRF14_BG and Subtype F1 Strains. <i>AIDS Research and Human Retroviruses</i> , 2015, 31, 488-495.	1.1	47
48	Big Events in Greece and HIV Infection Among People Who Inject Drugs. <i>Substance Use and Misuse</i> , 2015, 50, 825-838.	1.4	47
49	Primate immunodeficiency virus classification and nomenclature: Review. <i>Infection, Genetics and Evolution</i> , 2016, 46, 150-158.	2.3	47
50	Prevalence of resistance-associated mutations in newly diagnosed HIV-1 patients in Greece. <i>Virus Research</i> , 2005, 112, 115-122.	2.2	45
51	Antibodies against SARS-CoV-2 among health care workers in a country with low burden of COVID-19. <i>PLoS ONE</i> , 2020, 15, e0243025.	2.5	45
52	Increase in transmitted resistance to non-nucleoside reverse transcriptase inhibitors among newly diagnosed HIV-1 infections in Europe. <i>BMC Infectious Diseases</i> , 2014, 14, 407.	2.9	43
53	Dating the origin of hepatitis B virus reveals higher substitution rate and adaptation on the branch leading to F/H genotypes. <i>Molecular Phylogenetics and Evolution</i> , 2015, 93, 44-54.	2.7	43
54	HIV-1 molecular transmission clusters in nine European countries and Canada: association with demographic and clinical factors. <i>BMC Medicine</i> , 2019, 17, 4.	5.5	43

#	ARTICLE	IF	CITATIONS
55	Characterization of a Novel vpu -Harboring Simian Immunodeficiency Virus from a Dent's Mona Monkey (Cercopithecus mona denti). Journal of Virology, 2005, 79, 8560-8571.	3.4	42
56	Triple-Class Virologic Failure in HIV-Infected Patients Undergoing Antiretroviral Therapy for Up to 10 Years. Archives of Internal Medicine, 2010, 170, 410-419.	3.8	42
57	Molecular Analysis of the Full-Length Genome of HIV Type 1 Subtype I: Evidence of A/G/I Recombination. AIDS Research and Human Retroviruses, 1999, 15, 745-758.	1.1	41
58	Low-Dose IFN- γ Monotherapy in Treatment-Naïve Individuals with HIV-1 Infection: Evidence of Potent Suppression of Viral Replication. Journal of Interferon and Cytokine Research, 2001, 21, 861-869.	1.2	41
59	Trends in virological and clinical outcomes in individuals with HIV-1 infection and virological failure of drugs from three antiretroviral drug classes: a cohort study. Lancet Infectious Diseases, The, 2012, 12, 119-127.	9.1	41
60	Socially-Integrated Transdisciplinary HIV Prevention. AIDS and Behavior, 2014, 18, 1821-1834.	2.7	39
61	Integrating molecular epidemiology and social network analysis to study infectious diseases: Towards a socio-molecular era for public health. Infection, Genetics and Evolution, 2016, 46, 248-255.	2.3	37
62	National Income Inequality and Declining GDP Growth Rates Are Associated with Increases in HIV Diagnoses among People Who Inject Drugs in Europe: A Panel Data Analysis. PLoS ONE, 2015, 10, e0122367.	2.5	37
63	Enhanced HIV-1 surveillance using molecular epidemiology to study and monitor HIV-1 outbreaks among intravenous drug users (IDUs) in Athens and Bucharest. Infection, Genetics and Evolution, 2015, 35, 109-121.	2.3	36
64	Re-analysis of a human hepatitis B virus (HBV) isolate from an East African wild born Pan troglodytes schweinfurthii: Evidence for interspecies recombination between HBV infecting chimpanzee and human. Gene, 2005, 349, 165-171.	2.2	35
65	Monitoring quality and coverage of harm reduction services for people who use drugs: a consensus study. Harm Reduction Journal, 2017, 14, 19.	3.2	33
66	Immune-escape mutations and stop-codons in HBsAg develop in a large proportion of patients with chronic HBV infection exposed to anti-HBV drugs in Europe. BMC Infectious Diseases, 2018, 18, 251.	2.9	33
67	All-cause mortality in systemic rheumatic diseases under treatment compared with the general population, 2015-2019. RMD Open, 2021, 7, e001694.	3.8	33
68	Patterns of Transmitted HIV Drug Resistance in Europe Vary by Risk Group. PLoS ONE, 2014, 9, e94495.	2.5	32
69	Early hCG addition to rFSH for ovarian stimulation in IVF provides better results and the cDNA copies of the hCG receptor may be an indicator of successful stimulation. Reproductive Biology and Endocrinology, 2009, 7, 110.	3.3	31
70	Pre-exposure Prophylaxis (PrEP) Awareness and Use Within High HIV Transmission Networks. AIDS and Behavior, 2019, 23, 1893-1903.	2.7	31
71	Seroprevalence of Antibodies against SARS-CoV-2 among the Personnel and Students of the National and Kapodistrian University of Athens, Greece: A Preliminary Report. Life, 2020, 10, 214.	2.4	31
72	Effects of Social Distancing Measures during the First Epidemic Wave of Severe Acute Respiratory Syndrome Infection, Greece. Emerging Infectious Diseases, 2021, 27, 452-462.	4.3	31

#	ARTICLE	IF	CITATIONS
73	Analysis of the Evolutionary Relationships of HIV-1 and SIVcpz Sequences Using Bayesian Inference: Implications for the Origin of HIV-1. <i>Molecular Biology and Evolution</i> , 2003, 20, 1986-1996.	8.9	30
74	Genetic variants in immunoregulatory genes and risk for childhood lymphomas. <i>European Journal of Haematology</i> , 2009, 83, 334-342.	2.2	30
75	Upregulation of Human Endogenous Retroviruses in Bronchoalveolar Lavage Fluid of COVID-19 Patients. <i>Microbiology Spectrum</i> , 2021, 9, e0126021.	3.0	30
76	Hepatitis B Virus DNA Integration, Chronic Infections and Hepatocellular Carcinoma. <i>Microorganisms</i> , 2021, 9, 1787.	3.6	29
77	HIV-1 molecular epidemiology in the Balkans: a melting pot for high genetic diversity. <i>AIDS Reviews</i> , 2012, 14, 28-36.	1.0	29
78	Combined Analysis of the Prevalence of Drug-Resistant Hepatitis B Virus in Antiviral Therapy-Experienced Patients in Europe (CAPRE). <i>Journal of Infectious Diseases</i> , 2016, 213, 39-48.	4.0	28
79	Unravelling the history of hepatitis B virus genotypes A and D infection using a full-genome phylogenetic and phylogeographic approach. <i>ELife</i> , 2018, 7, .	6.0	28
80	Phylogenetic Reconstruction of a Known HIV-1 CRF04_cpx Transmission Network Using Maximum Likelihood and Bayesian Methods. <i>Journal of Molecular Evolution</i> , 2004, 59, 709-717.	1.8	26
81	Overview of HIV molecular epidemiology among people who inject drugs in Europe and Asia. <i>Infection, Genetics and Evolution</i> , 2016, 46, 256-268.	2.3	26
82	Prevalence of drug resistance among HIV-1 treatment-naïve patients in Greece during 2003-2015: Transmitted drug resistance is due to onward transmissions. <i>Infection, Genetics and Evolution</i> , 2017, 54, 183-191.	2.3	26
83	Outbreaks of SARS-CoV-2 in naturally infected mink farms: Impact, transmission dynamics, genetic patterns, and environmental contamination. <i>PLoS Pathogens</i> , 2021, 17, e1009883.	4.7	26
84	Molecular Characterization of a Recombinant HIV Type 1 Isolate (A/G/E?): Unidentified Regions May Be Derived from Parental Subtype E Sequences. <i>AIDS Research and Human Retroviruses</i> , 2000, 16, 845-855.	1.1	25
85	Network Characteristics of People Who Inject Drugs Within a New HIV Epidemic Following Austerity in Athens, Greece. <i>Journal of Acquired Immune Deficiency Syndromes (1999)</i> , 2015, 69, 499-508.	2.1	24
86	The application of HIV molecular epidemiology to public health. <i>Infection, Genetics and Evolution</i> , 2016, 46, 159-168.	2.3	24
87	HIV-1 Infection in Cyprus, the Eastern Mediterranean European Frontier: A Densely Sampled Transmission Dynamics Analysis from 1986 to 2012. <i>Scientific Reports</i> , 2018, 8, 1702.	3.3	24
88	EAES Recommendations for Recovery Plan in Minimally Invasive Surgery Amid COVID-19 Pandemic. <i>Surgical Endoscopy and Other Interventional Techniques</i> , 2021, 35, 1-17.	2.4	24
89	Prevalence patterns and genotypes of GB Virus C/hepatitis G virus among imprisoned intravenous drug users. , 1998, 56, 246-252.		23
90	Phylogenetic analysis of the full-length SARS-CoV sequences: Evidence for phylogenetic discordance in three genomic regions. <i>Journal of Medical Virology</i> , 2004, 74, 369-372.	5.0	23

#	ARTICLE	IF	CITATIONS
91	HIV epidemiology in Greece. <i>Future Microbiology</i> , 2008, 3, 507-516.	2.0	23
92	Development and Assessment of a Multiplex Real-Time PCR Assay for Quantification of Human Immunodeficiency Virus Type 1 DNA. <i>Journal of Clinical Microbiology</i> , 2009, 47, 2194-2199.	3.9	23
93	Development of a new ultra sensitive real-time PCR assay (ultra sensitive RTQ-PCR) for the quantification of HBV-DNA. <i>Virology Journal</i> , 2010, 7, 57.	3.4	23
94	Treatment-associated polymorphisms in protease are significantly associated with higher viral load and lower CD4 count in newly diagnosed drug-naïve HIV-1 infected patients. <i>Retrovirology</i> , 2012, 9, 81.	2.0	23
95	Rare occurrence of doravirine resistance-associated mutations in HIV-1-infected treatment-naïve patients. <i>Journal of Antimicrobial Chemotherapy</i> , 2019, 74, 614-617.	3.0	23
96	A Network Intervention to Locate Newly HIV Infected Persons Within MSM Networks in Chicago. <i>AIDS and Behavior</i> , 2019, 23, 15-20.	2.7	22
97	Genetic Features of HIV-1 Integrase Sub-Subtype A6 Predominant in Russia and Predicted Susceptibility to INSTIs. <i>Viruses</i> , 2020, 12, 838.	3.3	22
98	SlidingBayes: exploring recombination using a sliding window approach based on Bayesian phylogenetic inference. <i>Bioinformatics</i> , 2005, 21, 1274-1275.	4.1	21
99	Global and regional dispersal patterns of hepatitis B virus genotype E from and in Africa: A full-genome molecular analysis. <i>PLoS ONE</i> , 2020, 15, e0240375.	2.5	21
100	Re-analysis of human immunodeficiency virus type 1 isolates from Cyprus and Greece, initially designated as subtype I, reveals a unique complex A/G/H/K/? mosaic pattern. <i>Journal of General Virology</i> , 2001, 82, 575-580.	2.9	21
101	Detailed Molecular Epidemiologic Characterization of HIV-1 Infection in Bulgaria Reveals Broad Diversity and Evolving Phylodynamics. <i>PLoS ONE</i> , 2013, 8, e59666.	2.5	20
102	Molecular characterization of HIV-1 infection in Northwest Spain (2009–2013): Investigation of the subtype F outbreak. <i>Infection, Genetics and Evolution</i> , 2015, 30, 96-101.	2.3	20
103	The role of A-to-I RNA editing in infections by RNA viruses: Possible implications for SARS-CoV-2 infection. <i>Clinical Immunology</i> , 2021, 226, 108699.	3.2	20
104	Detailed Molecular Surveillance of the HIV-1 Outbreak Among People who Inject Drugs (PWID) in Athens During a Period of Four Years. <i>Current HIV Research</i> , 2018, 15, 396-404.	0.5	20
105	Alterations in cortisol and interleukin-6 secretion in patients with COVID-19 suggestive of neuroendocrine-immune adaptations. <i>Endocrine</i> , 2022, 75, 317-327.	2.3	20
106	Comparative evaluation of the QUANTIPLEX HIV-1 RNA 2.0 and 3.0 (bDNA) assays and the AMPLICOR HIV-1 MONITOR v1.5 test for the quantitation of human immunodeficiency virus type 1 RNA in plasma. <i>Journal of Virological Methods</i> , 2001, 91, 67-74.	2.1	19
107	HIV-1 genetic variants circulation in the North of Angola. <i>Infection, Genetics and Evolution</i> , 2005, 5, 231-237.	2.3	19
108	Molecular typing of the recently expanding subtype B HIV-1 epidemic in Romania: Evidence for local spread among MSMs in Bucharest area. <i>Infection, Genetics and Evolution</i> , 2012, 12, 1052-1057.	2.3	19

#	ARTICLE	IF	CITATIONS
109	Molecular Tracing of the Geographical Origin of Human Immunodeficiency Virus Type 1 Infection and Patterns of Epidemic Spread Among Migrants Who Inject Drugs in Athens. <i>Clinical Infectious Diseases</i> , 2017, 65, 2078-2084.	5.8	19
110	Transmission Dynamics of HIV-1 Drug Resistance among Treatment-Naïve Individuals in Greece: The Added Value of Molecular Epidemiology to Public Health. <i>Genes</i> , 2017, 8, 322.	2.4	19
111	Spatiotemporal Characteristics of the HIV-1 CRF02_AG/CRF63_02A1 Epidemic in Russia and Central Asia. <i>AIDS Research and Human Retroviruses</i> , 2018, 34, 415-420.	1.1	19
112	Temporal trends of transmitted HIV drug resistance in a multinational seroconversion cohort. <i>Aids</i> , 2018, 32, 161-169.	2.2	19
113	Genotypic Characterization of Human Immunodeficiency Virus Type 1 in Greece. <i>AIDS Research and Human Retroviruses</i> , 1998, 14, 685-690.	1.1	18
114	Quantitative Detection of the M204V Hepatitis B Virus Minor Variants by Amplification Refractory Mutation System Real-Time PCR Combined with Molecular Beacon Technology. <i>Journal of Clinical Microbiology</i> , 2009, 47, 2544-2550.	3.9	18
115	Prevalence of different HPV types and estimation of prognostic risk factors based on the linear array HPV genotyping test. <i>Journal of Medical Virology</i> , 2009, 81, 2059-2065.	5.0	18
116	Reducing HIV infection in people who inject drugs is impossible without targeting recently-infected subjects. <i>Aids</i> , 2016, 30, 2885-2890.	2.2	18
117	Evaluation of the limiting antigen avidity EIA (LAG) in people who inject drugs in Greece. <i>Epidemiology and Infection</i> , 2017, 145, 401-412.	2.1	18
118	Molecular Analysis of Human Immunodeficiency Virus Type 1 (HIV-1)â€“Infected Individuals in a Network-Based Intervention (Transmission Reduction Intervention Project): Phylogenetics Identify HIV-1â€“Infected Individuals With Social Links. <i>Journal of Infectious Diseases</i> , 2018, 218, 707-715.	4.0	18
119	Human Endogenous Retrovirus-K HML-2 integration within <i>RASGRF2</i> is associated with intravenous drug abuse and modulates transcription in a cell-line model. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 10434-10439.	7.1	18
120	Trends in COVID-19 Vaccination Intent, Determinants and Reasons for Vaccine Hesitancy: Results from Repeated Cross-Sectional Surveys in the Adult General Population of Greece during November 2020â€“June 2021. <i>Vaccines</i> , 2022, 10, 470.	4.4	18
121	HIV Transmission Patterns Among The Netherlands, Suriname, and The Netherlands Antilles: A Molecular Epidemiological Study. <i>AIDS Research and Human Retroviruses</i> , 2011, 27, 123-130.	1.1	17
122	Renal transplantation from hepatitis B surface antigen (<sc>HB</sc>sAg)â€“positive donors to <sc>HB</sc>sAgâ€“negative recipients: a case of postâ€“transplant fulminant hepatitis associated with an extensively mutated hepatitis B virus strain and review of the current literature. <i>Transplant Infectious Disease</i> , 2013, 15, 393-399.	1.7	17
123	Persistence of frequently transmitted drug-resistant HIV-1 variants can be explained by high viral replication capacity. <i>Retrovirology</i> , 2014, 11, 105.	2.0	17
124	Phylodynamics Helps to Evaluate the Impact of an HIV Prevention Intervention. <i>Viruses</i> , 2020, 12, 469.	3.3	17
125	Mutations associated with genotypic resistance to antiretroviral therapy in treatment naÃ±ve HIV-1 infected patients in Greece. <i>Virus Research</i> , 2002, 85, 109-115.	2.2	15
126	A contaminant-free assessment of Endogenous Retroviral RNA in human plasma. <i>Scientific Reports</i> , 2016, 6, 33598.	3.3	15

#	ARTICLE	IF	CITATIONS
127	Increased influenza vaccination rates in patients with autoimmune rheumatic diseases during the Covid-19 pandemic: a cross-sectional study. <i>Rheumatology International</i> , 2021, 41, 895-902.	3.0	15
128	MicroRNAs and HIV latency: a complex and promising relationship. <i>AIDS Reviews</i> , 2012, 14, 188-94.	1.0	15
129	HBV viremia in newborns of HBsAg(+) predominantly Caucasian HBeAg(â**) mothers. <i>Journal of Clinical Virology</i> , 2011, 50, 249-252.	3.1	14
130	Ultrasensitive Amplification Refractory Mutation System Real-Time PCR (ARMS RT-PCR) Assay for Detection of Minority Hepatitis B Virus-Resistant Strains in the Era of Personalized Medicine. <i>Journal of Clinical Microbiology</i> , 2013, 51, 2893-2900.	3.9	14
131	Differential Immunodominance Hierarchy of CD8⁺T-Cell Responses in HLA-B*27:05- and -B*27:02-Mediated Control of HIV-1 Infection. <i>Journal of Virology</i> , 2018, 92, .	3.4	14
132	Effect of immediate initiation of antiretroviral treatment on the risk of acquired HIV drug resistance. <i>Aids</i> , 2018, 32, 327-335.	2.2	13
133	Network Research Experiences in New York and Eastern Europe: Lessons for the Southern US in Understanding HIV Transmission Dynamics. <i>Current HIV/AIDS Reports</i> , 2018, 15, 283-292.	3.1	13
134	Calendar Time Trends in the Incidence and Prevalence of Triple-Class Virologic Failure in Antiretroviral Drug-Experienced People With HIV in Europe. <i>Journal of Acquired Immune Deficiency Syndromes</i> (1999), 2012, 59, 294-299.	2.1	13
135	Pre-Exposure Prophylaxis for HIV: Evidence and Perspectives. <i>Current Pharmaceutical Design</i> , 2017, 23, 2579-2591.	1.9	13
136	Modelling SARS-CoV-2 Binding Antibody Waning 8 Months after BNT162b2 Vaccination. <i>Vaccines</i> , 2022, 10, 285.	4.4	13
137	The interferon receptorâ€¹ promoter polymorphisms affect the outcome of Caucasians with <scp>HB</scp>eAgâ€¹negative chronic <scp>HBV</scp> infection. <i>Liver International</i> , 2015, 35, 2506-2513.	3.9	12
138	HBVâ€¹DNA levels predict overall mortality in HIV/HBV coinfecting individuals. <i>Journal of Medical Virology</i> , 2016, 88, 466-473.	5.0	12
139	SARS-CoV-2 Infection Is Asymptomatic in Nearly Half of Adults with Robust Anti-Spike Protein Receptor-Binding Domain Antibody Response. <i>Vaccines</i> , 2021, 9, 207.	4.4	12
140	Molecular characterization of a complex, recombinant human immunodeficiency virus type 1 (HIV-1) isolate (A/G/J/K/?): evidence to support the existence of a novel HIV-1 subtype. <i>Journal of General Virology</i> , 2001, 82, 2509-2514.	2.9	12
141	Recent Advances in Antiretroviral Agents: Potent Integrase Inhibitors. <i>Current Pharmaceutical Design</i> , 2017, 23, 2552-2567.	1.9	12
142	Molecular epidemiology of GB virus C/hepatitis G virus in Athens, Greece. <i>Journal of Medical Virology</i> , 2000, 61, 319-326.	5.0	11
143	Prevalence and genotypic distribution of TT virus in Athens, Greece. <i>Journal of Medical Virology</i> , 2001, 65, 423-429.	5.0	11
144	Assessment of phylogenetic sensitivity for reconstructing HIV-1 epidemiological relationships. <i>Virus Research</i> , 2012, 166, 54-60.	2.2	11

#	ARTICLE	IF	CITATIONS
145	Molecular investigation of HIV-1 cross-group transmissions during an outbreak among people who inject drugs (2011–2014) in Athens, Greece. <i>Infection, Genetics and Evolution</i> , 2018, 62, 11-16.	2.3	11
146	Effect of HIV Type 1 Subtype on Virological and Immunological Response to Combination Antiretroviral Therapy: Evidence for a More Rapid Viral Suppression for Subtype A Than Subtype B-Infected Greek Individuals. <i>AIDS Research and Human Retroviruses</i> , 2013, 29, 461-469.	1.1	10
147	Experiences of Stigma and Support Reported by Participants in a Network Intervention to Reduce HIV Transmission in Athens, Greece; Odessa, Ukraine; and Chicago, Illinois. <i>AIDS and Behavior</i> , 2019, 23, 1210-1224.	2.7	10
148	Understanding of Young Adolescents About HPV Infection: How Health Education Can Improve Vaccination Rate. <i>Journal of Cancer Education</i> , 2020, 35, 850-859.	1.3	10
149	Comparative Immunogenicity of BNT162b2 mRNA Vaccine with Natural SARS-CoV-2 Infection. <i>Vaccines</i> , 2021, 9, 1017.	4.4	10
150	Neuroaspis PLP10 α , ϕ , a nutritional formula rich in omega-3 and omega-6 fatty acids with antioxidant vitamins including gamma-tocopherol in early Parkinson's disease: A randomized, double-blind, placebo-controlled trial. <i>Clinical Neurology and Neurosurgery</i> , 2021, 210, 106954.	1.4	10
151	Cellular HIV-1 DNA Levels in Drug Sensitive Strains Are Equivalent to Those in Drug Resistant Strains in Newly-Diagnosed Patients in Europe. <i>PLoS ONE</i> , 2010, 5, e10976.	2.5	10
152	Blood Transcriptomes of Anti-SARS-CoV-2 Antibody-Positive Healthy Individuals Who Experienced Asymptomatic Versus Clinical Infection. <i>Frontiers in Immunology</i> , 2021, 12, 746203.	4.8	10
153	Clinical evaluation of an HIV-1 and HCV assay and demonstration of significant reduction of the HCV detection window before seroconversion. <i>Transfusion</i> , 2004, 44, 59-66.	1.6	9
154	Spatiotemporal Characteristics of the Largest HIV-1 CRF02_AG Outbreak in Spain: Evidence for Onward Transmissions. <i>Frontiers in Microbiology</i> , 2019, 10, 370.	3.5	9
155	Identifying, linking, and treating people who inject drugs and were recently infected with HIV in the context of a network-based intervention. <i>AIDS Care - Psychological and Socio-Medical Aspects of AIDS/HIV</i> , 2019, 31, 1376-1383.	1.2	9
156	Molecular Epidemiology Analysis of SARS-CoV-2 Strains Circulating in Romania during the First Months of the Pandemic. <i>Life</i> , 2020, 10, 152.	2.4	9
157	Prevalence and clinical implications of respiratory viruses in stable chronic obstructive pulmonary disease (COPD) and exacerbations: a systematic review and meta-analysis protocol. <i>BMJ Open</i> , 2020, 10, e035640.	1.9	9
158	Temporal Dominance of B.1.1.7 over B.1.354 SARS-CoV-2 Variant: A Hypothesis Based on Areas of Variant Co-Circulation. <i>Life</i> , 2021, 11, 375.	2.4	9
159	Discriminating Between Premigration and Postmigration HIV Acquisition Using Surveillance Data. <i>Journal of Acquired Immune Deficiency Syndromes (1999)</i> , 2021, 88, 117-124.	2.1	9
160	Prevalence patterns and genotypes of GB virus C/hepatitis G virus among imprisoned intravenous drug users. <i>Journal of Medical Virology</i> , 1998, 56, 246-52.	5.0	9
161	Short Communication: Dynamics of Drug-Resistant HIV-1 in Plasma and Peripheral Blood Cells in Patients during and after Enfuvirtide Therapy. <i>AIDS Research and Human Retroviruses</i> , 2007, 23, 1078-1082.	1.1	8
162	Analysing the evolutionary history of HCV: Puzzle of ancient phylogenetic discordance. <i>Infection, Genetics and Evolution</i> , 2007, 7, 354-360.	2.3	8

#	ARTICLE	IF	CITATIONS
163	Comparative evaluation of the performance of the Abbott RealTime HIV-1 assay for measurement of HIV-1 plasma viral load on genetically diverse samples from Greece. <i>Virology Journal</i> , 2011, 8, 10.	3.4	8
164	Impact of Interferon- γ Receptor-1 Promoter Polymorphisms on the Transcriptome of the Hepatitis B Virus-Associated Hepatocellular Carcinoma. <i>Frontiers in Immunology</i> , 2018, 9, 777.	4.8	8
165	Health and health needs of migrants in detention in Greece: shedding light to an unknown reality. <i>Globalization and Health</i> , 2019, 15, 4.	4.9	8
166	A Nationwide Study about the Dispersal Patterns of the Predominant HIV-1 Subtypes A1 and B in Greece: Inference of the Molecular Transmission Clusters. <i>Viruses</i> , 2020, 12, 1183.	3.3	8
167	Molecular Epidemiology of SARS-CoV-2 in Greece Reveals Low Rates of Onward Virus Transmission after Lifting of Travel Restrictions Based on Risk Assessment during Summer 2020. <i>MSphere</i> , 2021, 6, e0018021.	2.9	8
168	Identification of reverse transcriptase mutations associated with HIV-1 drug resistance mainly against non-nucleoside reverse transcriptase inhibitors in treatment-naïve patients. <i>Aids</i> , 1999, 13, 1276.	2.2	7
169	Predictors of CD4+ T-Cell Counts of HIV Type 1 "Infected Persons After Virologic Failure of All 3 Original Antiretroviral Drug Classes. <i>Journal of Infectious Diseases</i> , 2013, 207, 759-767.	4.0	7
170	An Innovative Study Design to Assess the Community Effect of Interventions to Mitigate HIV Epidemics Using Transmission-Chain Phylodynamics. <i>American Journal of Epidemiology</i> , 2018, 187, 2615-2622.	3.4	7
171	A hepatitis C outbreak preceded the HIV outbreak among persons who inject drugs in Athens, Greece: Insights from a mathematical modelling study. <i>Journal of Viral Hepatitis</i> , 2019, 26, 1311-1317.	2.0	7
172	SARS-CoV-2 Molecular Transmission Clusters and Containment Measures in Ten European Regions during the First Pandemic Wave. <i>Life</i> , 2021, 11, 219.	2.4	7
173	High prevalence of CRF02_AG and many minor resistance-related mutations at the protease gene among HIV-infected treatment-naïve immigrants in Madrid. <i>Aids</i> , 2003, 17, 1105-1107.	2.2	7
174	Molecular epidemiology of vertical human immunodeficiency virus type 1 transmission in Greece: evidence of non-B subtypes. <i>Journal of Human Virology</i> , 1999, 2, 339-43.	0.8	7
175	A Proposed Etiology for an Aberrant Response to Enteric Adenovirus Infection in Previously SARS-CoV-2-Infected Children With Acute Hepatitis. <i>Journal of the Pediatric Infectious Diseases Society</i> , 2022, 11, 352-353.	1.3	7
176	The Role of HIV-1 DNA as an Additional Marker of HIV-1 Infection. <i>Current HIV Research</i> , 2009, 7, 255-265.	0.5	6
177	Boosted Lopinavir " Versus Boosted Atazanavir "Containing Regimens and Immunologic, Virologic, and Clinical Outcomes: A Prospective Study of HIV-Infected Individuals in High-Income Countries. <i>Clinical Infectious Diseases</i> , 2015, 60, 1262-1268.	5.8	6
178	HIV-1 transmission networks across Cyprus (2010-2012). <i>PLoS ONE</i> , 2018, 13, e0195660.	2.5	6
179	Effectiveness of Transmitted Drug Resistance Testing Before Initiation of Antiretroviral Therapy in HIV-Positive Individuals. <i>Journal of Acquired Immune Deficiency Syndromes (1999)</i> , 2019, 82, 314-320.	2.1	6
180	The Effect of Health Education on Adolescents' Awareness of HPV Infections and Attitudes towards HPV Vaccination in Greece. <i>International Journal of Environmental Research and Public Health</i> , 2022, 19, 503.	2.6	6

#	ARTICLE	IF	CITATIONS
181	Role of genetic polymorphisms in the progression of liver fibrosis in chronic hepatitis C virus infection. <i>Liver International</i> , 2008, 28, 764-766.	3.9	5
182	Emergence of an NNRTI Resistance Mutation Y181C in an HIV-Infected NNRTI-Naive Patient. <i>AIDS Research and Human Retroviruses</i> , 2008, 24, 413-415.	1.1	5
183	High prevalence of the UGT1A1*28 variant in HIV-infected individuals in Greece. <i>International Journal of STD and AIDS</i> , 2014, 25, 860-865.	1.1	5
184	Hyperbilirubinemia in atazanavir treated HIV-infected patients: the impact of the UGT1A1*28 allele. <i>Pharmacogenomics and Personalized Medicine</i> , 2017, Volume 10, 205-208.	0.7	5
185	HCV Phylogeography of the General Population and High-Risk Groups in Cyprus Identifies the Island as a Global Sink for and Source of Infection. <i>Scientific Reports</i> , 2019, 9, 10077.	3.3	5
186	Allele Frequency of the CCR5 Mutant Chemokine Receptor in Greek Caucasians. <i>Journal of Acquired Immune Deficiency Syndromes</i> , 1998, 17, 181-182.	0.3	4
187	Re-analysis of 34 full-length HIV-1 intersubtype recombinant sequences. <i>Infection, Genetics and Evolution</i> , 2005, 5, 225-229.	2.3	4
188	Monitoring HIV drug resistance in treatment-naive individuals: molecular indicators, epidemiology and clinical implications. <i>HIV Therapy</i> , 2009, 3, 369-390.	0.6	4
189	Pockets of HIV Non-infection Within Highly-Infected Risk Networks in Athens, Greece. <i>Frontiers in Microbiology</i> , 2018, 9, 1825.	3.5	4
190	Spatiotemporal Distribution and Genetic Characterization of Measles Strains Circulating in Greece during the 2017–2018 Outbreak. <i>Viruses</i> , 2020, 12, 1166.	3.3	4
191	Sexual Mixing and HIV Transmission Potential Among Greek Men Who have Sex with Men: Results from SOPHOCLES. <i>AIDS and Behavior</i> , 2021, 25, 1935-1945.	2.7	4
192	Validation of molecular clock inferred HIV infection ages: Evidence for accurate estimation of infection dates. <i>Infection, Genetics and Evolution</i> , 2021, 91, 104799.	2.3	4
193	People with high HIV viral load within risk networks: who are these people and who refers them best?. <i>Journal of Infection in Developing Countries</i> , 2019, 13, 103S-110S.	1.2	4
194	Ongoing HIV transmission following a large outbreak among people who inject drugs in Athens, Greece (2014–20). <i>Addiction</i> , 2022, 117, 1670-1682.	3.3	4
195	Vaccination Hesitancy among Health-Care-Workers in Academic Hospitals Is Associated with a 12-Fold Increase in the Risk of COVID-19 Infection: A Nine-Month Greek Cohort Study. <i>Viruses</i> , 2022, 14, 26.	3.3	4
196	Analysis of complex HIV-1 intersubtype recombinants using a Bayesian scanning method. <i>Infection, Genetics and Evolution</i> , 2005, 5, 219-224.	2.3	3
197	HIV-1 epidemic in Russia: an evolutionary epidemiology analysis. <i>Lancet, The</i> , 2014, 383, S71.	13.7	3
198	HCV dispersal patterns among intravenous drug users (IDUs) in Athens metropolitan area. <i>Infection, Genetics and Evolution</i> , 2016, 45, 415-419.	2.3	3

#	ARTICLE	IF	CITATIONS
199	Near Full-length Genomic Sequencing and Molecular Analysis of HIV-Infected Individuals in a Network-based Intervention (TRIP) in Athens, Greece: Evidence that Transmissions Occur More Frequently from those with High HIV-RNA. <i>Current HIV Research</i> , 2019, 16, 345-353.	0.5	3
200	Automated, phylogeny-based genotype delimitation of the Hepatitis Viruses HBV and HCV. <i>PeerJ</i> , 2019, 7, e7754.	2.0	3
201	Effects of low molecular weight heparin and fondaparinux on mortality, hemorrhagic and thrombotic complications in COVID-19 patients. <i>Therapeutic Advances in Neurological Disorders</i> , 2022, 15, 175628642210994.	3.5	3
202	Genetic Evolution of Human Immunodeficiency Virus Type 1 in Two Spouses Responding Successfully to Highly Active Antiretroviral Therapy. <i>AIDS Research and Human Retroviruses</i> , 2003, 19, 65-71.	1.1	2
203	Correlation between patterns of HIV-1 drug resistance and drug administration in antiretroviral experienced patients in Greece during 1999–2006. <i>Journal of the International AIDS Society</i> , 2008, 11, P186.	3.0	2
204	Comments on “The distribution of HIV-1 recombination breakpoints” by Jun Fan, Matteo Negroni, David L. Robertson. <i>Infection, Genetics and Evolution</i> , 2008, 8, 527-528.	2.3	2
205	HIV/HBV co-infection and rate of antiretroviral treatment change after highly active antiretroviral treatment initiation in a cohort of HIV-infected patients in Greece. <i>International Journal of STD and AIDS</i> , 2010, 21, 702-707.	1.1	2
206	Re-analysis of unassigned hepatitis C virus (HCV) strain CYHCV025: Evidence of a highly divergent lineage within genotype 1. <i>Molecular Phylogenetics and Evolution</i> , 2017, 109, 362-366.	2.7	2
207	Investigation of Human Immunodeficiency Virus Outbreaks Among People Who Inject Drugs: Timely Diagnosis and Molecular Surveillance are Crucial. <i>Journal of Infectious Diseases</i> , 2017, 216, 1049-1050.	4.0	2
208	Global molecular epidemiology of HIV-1: the chameleon challenge. <i>Lancet Infectious Diseases</i> , The, 2019, 19, 114-115.	9.1	2
209	Earlier treatment initiation is associated with a decreased number of HIV-1 subtype A1 transmissions in Greece. <i>Sexually Transmitted Infections</i> , 2021, 97, 232-237.	1.9	2
210	Tracking missed opportunities for an early HIV diagnosis in a population of people living with HIV with known time of infection. <i>Sexually Transmitted Infections</i> , 2021, , sextrans-2020-054697.	1.9	2
211	Assessment of HIV testing recommendations in Greek specialty guidelines: A missed opportunity and room for improvement for recommending testing. <i>AIDS Care - Psychological and Socio-Medical Aspects of AIDS/HIV</i> , 2021, 33, 1312-1315.	1.2	2
212	Integrated molecular and affiliation network analysis: Core-periphery social clustering is associated with HIV transmission patterns. <i>Social Networks</i> , 2022, 68, 107-117.	2.1	2
213	Authors’ reply: HIV-1 outbreak among injecting drug users in Greece, 2011: a preliminary report. <i>Eurosurveillance</i> , 2011, 16, .	7.0	2
214	Presenting a conceptual framework for an HIV prevention and care continuum and assessing the feasibility of empirical measurement in Estonia: A case study. <i>PLoS ONE</i> , 2020, 15, e0240224.	2.5	2
215	Dating the Origin and Estimating the Transmission Rates of the Major HIV-1 Clusters in Greece: Evidence about the Earliest Subtype A1 Epidemic in Europe. <i>Viruses</i> , 2022, 14, 101.	3.3	2
216	Modeling the impact of interventions during an outbreak of HIV infection among people who inject drugs in 2012–2013 in Athens, Greece.. <i>Drug and Alcohol Dependence</i> , 2022, 234, 109396.	3.2	2

#	ARTICLE	IF	CITATIONS
217	THU-427-A community-based fast-track seek-test-treat program to enhance diagnosis and linkage to care for hepatitis C infection among people who inject drugs in Athens, Greece (ARISTOTLE HCV-HIV) Tj ETQq1 1 0.784317 rgBT / @Overlock 10		
218	Classification, Genetic Diversity and Global Distribution of Hepatitis C Virus (HCV) Genotypes and Subtypes. , 2021, , 55-69.		1
219	Sociometric Risk Network Structure, HIV Prevalence, and Drug Injection-Related Norms among People Who Inject Drugs (PWID) in Athens, Greece. Substance Use and Misuse, 2021, 56, 1190-1201.	1.4	1
220	Pre-exposure Prophylaxis Persistence Among Greek Sexual Minority Men: Results from PrEP for Greece (P4G) Study. AIDS and Behavior, 2021, , 1.	2.7	1
221	Identifying Country-Level Risk Factors for the Spread of COVID-19 in Europe Using Machine Learning. Viruses, 2022, 14, 625.	3.3	1
222	Reanalysis of the HIV-1 Circulating Recombinant Form A/E (CRF01_AE): Evidence of A/E/G Recombination. Journal of Acquired Immune Deficiency Syndromes (1999), 2002, 30, 124-129.	2.1	0
223	409 dating the HCV3a epidemic in greece using molecular clock approaches. Journal of Hepatology, 2006, 44, S154.	3.7	0
224	[479] PERINATAL HBV VIREMIA IN CHILDREN BORN TO HBsAg(+)/HBeAg(â”) MOTHERS. Journal of Hepatology, 2007, 46, S182.	3.7	0
225	The phylogenetic information profile of HIV-1 and the degradation effect of recombination. Infection, Genetics and Evolution, 2008, 8, 139-145.	2.3	0
226	570 CLINICAL EVALUATION OF THE ARMS RT-PCR FOR THE QUANTITATIVE DETECTION OF M204V VARIANTS IN CHRONIC HBV PATIENTS. Journal of Hepatology, 2008, 48, S214.	3.7	0
227	573 DEVELOPMENT OF A HIGHLY SENSITIVE SYSTEM FOR THE DETECTION AND QUANTIFICATION OF ALL MAJOR HBV RESISTANCE MUTATIONS BASED ON ARMS RT-PCR WITH MOLECULAR BEACON. Journal of Hepatology, 2009, 50, S211.	3.7	0
228	Successful virological outcome in an HIV-infected individual with a three-class resistant variant and an insertion in the protease genome with a Tipranavir based regimen. Irish Journal of Medical Science, 2010, 179, 305-307.	1.5	0
229	High prevalence of the UGT1A1*28 variant in HIV-infected individuals in Greece. Journal of the International AIDS Society, 2010, 13, P146-P146.	3.0	0
230	1117 DETECTION OF MINORITY HBV RESISTANT STRAINS IN HBV TREATMENT NAIVE PATIENTS USING A HIGHLY SENSITIVE ASSAY BASED ON ARMS RT-PCR WITH MOLECULAR BEACON. Journal of Hepatology, 2011, 54, S442.	3.7	0
231	Appearance of a Single Amino Acid Insertion at Position 33 in HIV Type 1 Protease Under a Lopinavir-Containing Regimen, Associated with Reduced Protease Inhibitor Susceptibility. AIDS Research and Human Retroviruses, 2011, 27, 1223-1229.	1.1	0
232	Patterns of drug resistance among newly diagnosed HIV-1 infected patients in Greece during the last decade: the crucial role of transmission networks. Journal of the International AIDS Society, 2014, 17, 19742.	3.0	0
233	P0537 : Dating the origin of HBV using full-length sequences. Journal of Hepatology, 2015, 62, S516.	3.7	0
234	P0538 : Tracing HBV genetic history in the arctic: Evidence for a central asian HBV origin in Greenland. Journal of Hepatology, 2015, 62, S517.	3.7	0

#	ARTICLE	IF	CITATIONS
235	P0547 : Phylogeography of global HBV genotype D reveals the central role of central Asia in disseminating the epidemic. Journal of Hepatology, 2015, 62, S520.	3.7	0
236	P0567 : Global patterns of HBV genotype D epidemic dispersal. Journal of Hepatology, 2015, 62, S527-S528.	3.7	0
237	Editorial. Infection, Genetics and Evolution, 2016, 46, 149.	2.3	0
238	Immune-Escape Mutations and Stop-Codons in HBsAg Circulates in a Relevant Proportion of Patients with Chronic HBV Infection Exposed to Anti-HBV Drugs in Europe: Implications for HBV Transmission in the Setting of Vaccination and Disease Progression. Journal of Hepatology, 2016, 64, S367-S368.	3.7	0
239	O14.3â€œ...How war and risky sexual behaviours shape the ukrainian hiv epidemic: a phylogeographic analysis. , 2017, , .		0
240	Factors associated with failure to link people who inject drugs to HCV care and treatment: results from a community-based seek-test-treat program in Athens, Greece (ARISTOTLE HCV-HIV). Journal of Hepatology, 2020, 73, S830.	3.7	0
241	An evolving genetic tapestry of HIV-1 recombinants. Lancet HIV,the, 2020, 7, e733-e734.	4.7	0
242	Hepatitis C Virus Origin. , 2021, , 45-53.		0
243	Development of a novel real-time PCR assay for the quantitation of HBV DNA. Journal of Hepatology, 2001, 34, 129.	3.7	0
244	A Network Approach to Determine Optimization of PrEP Uptake in Athens, Greece. AIDS and Behavior, 2022, , 1.	2.7	0
245	Antibodies against SARS-CoV-2 among health care workers in a country with low burden of COVID-19. , 2020, 15, e0243025.		0
246	Antibodies against SARS-CoV-2 among health care workers in a country with low burden of COVID-19. , 2020, 15, e0243025.		0
247	Antibodies against SARS-CoV-2 among health care workers in a country with low burden of COVID-19. , 2020, 15, e0243025.		0
248	Antibodies against SARS-CoV-2 among health care workers in a country with low burden of COVID-19. , 2020, 15, e0243025.		0