Evangelia-Georgia Kostaki

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3997883/publications.pdf

Version: 2024-02-01



#	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. Infection, Genetics and Evolution, 2020, 79, 104212.	1.0	544
2	Phylogenetic Analysis of SARS-CoV-2 Data Is Difficult. Molecular Biology and Evolution, 2021, 38, 1777-1791.	3.5	119
3	Molecular epidemiology of HIV-1 infection in Europe: An overview. Infection, Genetics and Evolution, 2016, 46, 180-189.	1.0	68
4	Immunomodulatory Function of HBeAg Related to Short-Sighted Evolution, Transmissibility, and Clinical Manifestation of Hepatitis B Virus. Frontiers in Microbiology, 2018, 9, 2521.	1.5	62
5	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. Science of the Total Environment, 2021, 768, 144578.	3.9	59
6	Dating the origin of hepatitis B virus reveals higher substitution rate and adaptation on the branch leading to F/H genotypes. Molecular Phylogenetics and Evolution, 2015, 93, 44-54.	1.2	43
7	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.	1.1	31
8	Unravelling the history of hepatitis B virus genotypes A and D infection using a full-genome phylogenetic and phylogeographic approach. ELife, 2018, 7, .	2.8	28
9	Overview of HIV molecular epidemiology among people who inject drugs in Europe and Asia. Infection, Genetics and Evolution, 2016, 46, 256-268.	1.0	26
10	Prevalence of drug resistance among HIV-1 treatment-naive patients in Greece during 2003–2015: Transmitted drug resistance is due to onward transmissions. Infection, Genetics and Evolution, 2017, 54, 183-191.	1.0	26
11	Clobal and regional dispersal patterns of hepatitis B virus genotype E from and in Africa: A full-genome molecular analysis. PLoS ONE, 2020, 15, e0240375.	1.1	21
12	Molecular characterization of HIV-1 infection in Northwest Spain (2009–2013): Investigation of the subtype F outbreak. Infection, Genetics and Evolution, 2015, 30, 96-101.	1.0	20
13	Detailed Molecular Surveillance of the HIV-1 Outbreak Among People who Inject Drugs (PWID) in Athens During a Period of Four Years. Current HIV Research, 2018, 15, 396-404.	0.2	20
14	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. Clinical Infectious Diseases, 2017, 65, 2078-2084.	2.9	19
15	Transmission Dynamics of HIV-1 Drug Resistance among Treatment-NaÃ⁻ve Individuals in Greece: The Added Value of Molecular Epidemiology to Public Health. Genes, 2017, 8, 322.	1.0	19
16	Spatiotemporal Characteristics of the HIV-1 CRF02_AG/CRF63_02A1 Epidemic in Russia and Central Asia. AIDS Research and Human Retroviruses, 2018, 34, 415-420.	0.5	19
17	Molecular Analysis of Human Immunodeficiency Virus Type 1 (HIV-1)–Infected Individuals in a Network-Based Intervention (Transmission Reduction Intervention Project): Phylogenetics Identify HIV-1–Infected Individuals With Social Links. Journal of Infectious Diseases, 2018, 218, 707-715.	1.9	18
18	Network Research Experiences in New York and Eastern Europe: Lessons for the Southern US in Understanding HIV Transmission Dynamics. Current HIV/AIDS Reports, 2018, 15, 283-292.	1.1	13

#	Article	IF	CITATIONS
19	SARS-CoV-2 Infection Is Asymptomatic in Nearly Half of Adults with Robust Anti-Spike Protein Receptor-Binding Domain Antibody Response. Vaccines, 2021, 9, 207.	2.1	12
20	Molecular investigation of HIV-1 cross-group transmissions during an outbreak among people who inject drugs (2011–2014) in Athens, Greece. Infection, Genetics and Evolution, 2018, 62, 11-16.	1.0	11
21	Neuroaspis PLP10â"¢, 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. Clinical Neurology and Neurosurgery, 2021, 210, 106954.	0.6	10
22	Spatiotemporal Characteristics of the Largest HIV-1 CRF02_AG Outbreak in Spain: Evidence for Onward Transmissions. Frontiers in Microbiology, 2019, 10, 370.	1.5	9
23	Temporal Dominance of B.1.1.7 over B.1.354 SARS-CoV-2 Variant: A Hypothesis Based on Areas of Variant Co-Circulation. Life, 2021, 11, 375.	1.1	9
24	A Nationwide Study about the Dispersal Patterns of the Predominant HIV-1 Subtypes A1 and B in Greece: Inference of the Molecular Transmission Clusters. Viruses, 2020, 12, 1183.	1.5	8
25	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. MSphere, 2021, 6, e0018021.	1.3	8
26	SARS-CoV-2 Molecular Transmission Clusters and Containment Measures in Ten European Regions during the First Pandemic Wave. Life, 2021, 11, 219.	1.1	7
27	HIV-1 transmission networks across Cyprus (2010-2012). PLoS ONE, 2018, 13, e0195660.	1.1	6
28	Pockets of HIV Non-infection Within Highly-Infected Risk Networks in Athens, Greece. Frontiers in Microbiology, 2018, 9, 1825.	1.5	4
29	Spatiotemporal Distribution and Genetic Characterization of Measles Strains Circulating in Greece during the 2017–2018 Outbreak. Viruses, 2020, 12, 1166.	1.5	4
30	Validation of molecular clock inferred HIV infection ages: Evidence for accurate estimation of infection dates. Infection, Genetics and Evolution, 2021, 91, 104799.	1.0	4
31	Ongoing HIV transmission following a large outbreak among people who inject drugs in Athens, Greece (2014–20). Addiction, 2022, 117, 1670-1682.	1.7	4
32	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. Viruses, 2022, 14, 26.	1.5	4
33	Near Full-length Genomic Sequencing and Molecular Analysis of HIVInfected Individuals in a Network-based Intervention (TRIP) in Athens, Greece: Evidence that Transmissions Occur More Frequently from those with High HIV-RNA. Current HIV Research, 2019, 16, 345-353.	0.2	3
34	Automated, phylogeny-based genotype delimitation of the Hepatitis Viruses HBV and HCV. PeerJ, 2019, 7, e7754.	0.9	3
35	Earlier treatment initiation is associated with a decreased number of HIV-1 subtype A1 transmissions in Greece. Sexually Transmitted Infections, 2021, 97, 232-237.	0.8	2
36	Tracking missed opportunities for an early HIV diagnosis in a population of people living with HIV with known time of infection. Sexually Transmitted Infections, 2021, , sextrans-2020-054697.	0.8	2

#	Article	IF	CITATIONS
37	Assessment of HIV testing recommendations in Greek specialty guidelines: A missed opportunity and room for improvement for recommending testing. AIDS Care - Psychological and Socio-Medical Aspects of AIDS/HIV, 2021, 33, 1312-1315.	0.6	2
38	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. Viruses, 2022, 14, 101.	1.5	2
39	Classification, Genetic Diversity and Global Distribution of Hepatitis C Virus (HCV) Genotypes and Subtypes. , 2021, , 55-69.		1
40	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.	0.7	1
41	Women in the European Virus Bioinformatics Center. Viruses, 2022, 14, 1522.	1.5	1
42	SAT-173-Global patterns of HBV genotype E epidemic dispersal from Africa determined using a full-genome phylogenetic and phylogeographic approach. Journal of Hepatology, 2019, 70, e706.	1.8	0
43	An evolving genetic tapestry of HIV-1 recombinants. Lancet HIV,the, 2020, 7, e733-e734.	2.1	0