Angeliki Melidou

List of Publications by Year in descending order

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23 23 23 1487 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	Geographical and temporal distribution of SARS-CoV-2 clades in the WHO European Region, January to June 2020. Eurosurveillance, 2020, 25, .	3.9	186
2	Very little influenza in the WHO European Region during the 2020/21 season, weeks 40 2020 to 8 2021. Eurosurveillance, 2021, 26, .	3.9	74
3	Dominant influenza A(H3N2) and B/Yamagata virus circulation in EU/EEA, 2016/17 and 2017/18 seasons, respectively. Eurosurveillance, 2018, 23, .	3.9	56
4	Molecular and phylogenetic analysis of the haemagglutinin gene of pandemic influenza H1N1 2009 viruses associated with severe and fatal infections. Virus Research, 2010, 151, 192-199.	1,1	52
5	Proficiency Testing of Virus Diagnostics Based on Bioinformatics Analysis of Simulated <i>In Silico</i> Ii> High-Throughput Sequencing Data Sets. Journal of Clinical Microbiology, 2019, 57, .	1.8	34
6	Paediatric Virology: A rapidly increasing educational challenge. Experimental and Therapeutic Medicine, 2017, 13, 364-377.	0.8	31
7	Virological surveillance of influenza viruses in the WHO European Region in 2019/20 – impact of the COVID-19 pandemic. Eurosurveillance, 2020, 25, .	3.9	24
8	Influenza returns with a season dominated by clade 3C.2a1b.2a.2 A(H3N2) viruses, WHO European Region, 2021/22. Eurosurveillance, 2022, 27, .	3.9	20
9	Paediatric Virology and its interaction between basic science and clinical practice (Review). International Journal of Molecular Medicine, 2018, 41, 1165-1176.	1.8	19
10	MicroRNA-related polymorphisms in pseudoexfoliation syndrome, pseudoexfoliative glaucoma, and primary open-angle glaucoma. Ophthalmic Genetics, 2018, 39, 603-609.	0.5	18
11	A PCR-based NGS protocol for whole genome sequencing of West Nile virus lineage 2 directly from biological specimens. Molecular and Cellular Probes, 2019, 46, 101412.	0.9	17
12	A paediatric influenza update 100 years after the Skyros island Spanish flu outbreak. Experimental and Therapeutic Medicine, 2019, 17, 4327-4336.	0.8	14
13	Influenza vaccine effectiveness against laboratory confirmed influenza in Greece during the 2013–2014 season: A test-negative study. Vaccine, 2015, 33, 367-373.	1.7	13
14	Isolation and whole-genome sequencing of a Crimean-Congo hemorrhagic fever virus strain, Greece. Ticks and Tick-borne Diseases, 2018, 9, 788-791.	1.1	13
15	The possible role of oral microbiome in autoimmunity. International Journal of Women's Dermatology, 2020, 6, 357-364.	1.1	13
16	Influenza vaccine effectiveness in preventing hospitalizations with laboratory-confirmed influenza in Greece during the 2014-2015 season: A test-negative study. Journal of Medical Virology, 2016, 88, 1896-1904.	2.5	11
17	Genetic analysis of post-pandemic 2010-2011 influenza A(H1N1)pdm09 hemagglutinin virus variants that caused mild, severe, and fatal infections in Northern Greece. Journal of Medical Virology, 2015, 87, 57-67.	2.5	10
18	Detection of the tetM resistance determinant among phenotypically sensitive Ureaplasma species by a novel real-time PCR method. Diagnostic Microbiology and Infectious Disease, 2015, 81, 85-88.	0.8	9

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#	Article	IF	CITATIONS
19	Molecular and phylogenetic analysis and vaccine strain match of human influenza A(H3N2) viruses isolated in Northern Greece between 2004 and 2008. Virus Research, 2009, 145, 220-226.	1.1	8
20	The role of oral microbiome in pemphigus vulgaris. Archives of Microbiology, 2021, 203, 2237-2247.	1.0	6
21	Application of 16S rRNA next generation sequencing in ticks in Greece. Heliyon, 2020, 6, e04542.	1.4	3
22	Genetic characterization of livestock-associated methicillin-resistant Staphylococcus aureus isolated in Greece. Brazilian Journal of Microbiology, 2021, 52, 2091-2096.	0.8	2
23	Congenital rubella and hyperthyroidism: A case report. Journal of Pediatric Infectious Diseases, 2015, 03, 141-144.	0.1	0