Maja Stanojevic

List of Publications by Year in descending order

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Μαία Stanoifuic

#	Article	IF	CITATIONS
1	Spike-directed vaccination elicits robust spike-specific T-cell response, including to mutant strains. Cytotherapy, 2022, 24, 10-15.	0.3	6
2	Tumor-associated antigen–specific T cells with nivolumab are safe and persist in vivo in relapsed/refractory Hodgkin lymphoma. Blood Advances, 2022, 6, 473-485.	2.5	11
3	The generation and application of antigen-specific TÂcell therapies for cancer and viral-associated disease. Molecular Therapy, 2022, 30, 2130-2152.	3.7	19
4	Outcome of donor-derived TAA-T cell therapy in patients with high-risk or relapsed acute leukemia post allogeneic BMT. Blood Advances, 2022, 6, 2520-2534.	2.5	19
5	Identification of novel HLA-restricted preferentially expressed antigen in melanoma peptides to facilitate off-the-shelf tumor-associated antigen-specific T-cell therapies. Cytotherapy, 2021, 23, 694-703.	0.3	7
6	A Laboratory-Based Surveillance Study of Invasive Neisseria meningitidis, Streptococcus pneumoniae, and Haemophilus influenzae Diseases in a Serbian Pediatric Population—Implications for Vaccination. Diagnostics, 2021, 11, 1059.	1.3	2
7	Colonization with Multidrug-Resistant Bacteria in the First Week of Life among Hospitalized Preterm Neonates in Serbia: Risk Factors and Outcomes. Microorganisms, 2021, 9, 2613.	1.6	10
8	Depicting the RNA Virome of Hematophagous Arthropods from Belgrade, Serbia. Viruses, 2020, 12, 975.	1.5	19
9	Tumor Associated Antigen Specific T Cells Given in Combination with Nivolumab for the Treatment of Hodgkin Lymphoma. Blood, 2020, 136, 18-18.	0.6	1
10	Castleman's disease associated with mixed connective tissue disorder and cerebral ischaemia and vasculitis: A rare case and a diagnostic challenge for an infectologist. Vojnosanitetski Pregled, 2020, 77, 872-877.	0.1	0
11	Exploring Evolutionary and Transmission Dynamics of HIV Epidemic in Serbia: Bridging Socio-Demographic With Phylogenetic Approach. Frontiers in Microbiology, 2019, 10, 287.	1.5	15
12	Molecular characterization of Dobrava-Belgrade hantavirus in Serbia, 2007–2011. Journal of Infection and Public Health, 2019, 12, 645-649.	1.9	3
13	Insight into diversity of bacteria belonging to the order Rickettsiales in 9 arthropods species collected in Serbia. Scientific Reports, 2019, 9, 18680.	1.6	23
14	Hepatitis B Outbreak Among Men Who Have Sex with Men in the Autonomous Province of Vojvodina, Serbia. LGBT Health, 2018, 5, 91-93.	1.8	3
15	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.	1.3	33
16	Forensic application of phylogenetic analyses – Exploration of suspected HIV-1 transmission case. Forensic Science International: Genetics, 2017, 27, 100-105.	1.6	18
17	High Frequency of Human Leukocyte Antigen-B*57:01 Allele Carriers among HIV-Infected Patients in Serbia. Intervirology, 2017, 60, 43-47.	1.2	0
18	Intravenous drug use – an independent predictor for HCV genotypes 3 and 4 infection among HIV/HCV co-infected patients. Archives of Medical Science, 2017, 3, 652-658.	0.4	13

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19	Locally advanced rectal cancers with simultaneous occurrence of KRAS mutation and high VEGF expression show invasive characteristics. Pathology Research and Practice, 2016, 212, 598-603.	1.0	12
20	Rubella immune status of neonates – a window towards seroprevalence among childbearing women. BMC Public Health, 2016, 16, 838.	1.2	3
21	Isolation and Molecular Detection of Bovine Parainfluenza Virus Type 3 in Cattle in Serbia. Acta Veterinaria, 2016, 66, 509-519.	0.2	1
22	The global spread of HIV-1 subtype B epidemic. Infection, Genetics and Evolution, 2016, 46, 169-179.	1.0	60
23	The influence of host factors and sequence variability of the p7 region on the response to pegylated interferon/ribavirin therapy for chronic hepatitis C genotype 1b in patients from Serbia. Archives of Virology, 2016, 161, 1189-1198.	0.9	2
24	Transmission of HIV Drug Resistance and the Predicted Effect on Current First-line Regimens in Europe. Clinical Infectious Diseases, 2016, 62, 655-663.	2.9	135
25	Combined Analysis of the Prevalence of Drug-Resistant Hepatitis B Virus in Antiviral Therapy–Experienced Patients in Europe (CAPRE). Journal of Infectious Diseases, 2016, 213, 39-48.	1.9	28
26	Molecular Detection of PCV2 And PPV in Pigs in Republic of Srpska, Bosnia and Herzegovina. Acta Veterinaria, 2016, 66, 51-60.	0.2	5
27	Geographic and Temporal Trends in the Molecular Epidemiology and Genetic Mechanisms of Transmitted HIV-1 Drug Resistance: An Individual-Patient- and Sequence-Level Meta-Analysis. PLoS Medicine, 2015, 12, e1001810.	3.9	188
28	Molecular characterization of macrolide resistant Streptococcus pyogenes isolates from pharyngitis patients in Serbia. Infection, Genetics and Evolution, 2015, 33, 246-252.	1.0	9
29	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. Journal of Infectious Diseases, 2015, 211, 1735-1744.	1.9	62
30	Natural Products as Promising Therapeutics for Treatment of Influenza Disease. Current Pharmaceutical Design, 2015, 21, 5573-5588.	0.9	27
31	Trends and Predictors of Transmitted Drug Resistance (TDR) and Clusters with TDR in a Local Belgian HIV-1 Epidemic. PLoS ONE, 2014, 9, e101738.	1.1	36
32	Ten Years Survey of Primary HIV-1 Resistance in Serbia: The Occurrence of Multiclass Resistance. AIDS Research and Human Retroviruses, 2014, 30, 634-641.	0.5	21
33	The Role and Therapeutic Potential of Autophagy Modulation in Controlling Virusâ€Induced Cell Death. Medicinal Research Reviews, 2014, 34, 744-767.	5.0	12
34	Evidence of recombination in Tula virus strains from Serbia. Infection, Genetics and Evolution, 2014, 21, 472-478.	1.0	19
35	Increase in transmitted resistance to non-nucleoside reverse transcriptase inhibitors among newly diagnosed HIV-1 infections in Europe. BMC Infectious Diseases, 2014, 14, 407.	1.3	43
36	Patterns of Transmitted HIV Drug Resistance in Europe Vary by Risk Group. PLoS ONE, 2014, 9, e94495.	1.1	32

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37	Distribution of macrolide-resistant genes among isolates of macrolideresistant Streptococcus pyogenes and Streptococcus pneumoniae in Serbia. Archives of Biological Sciences, 2014, 66, 93-98.	0.2	5
38	Comparative phylogenetic analysis of Dobrava-Belgrade virus L and S genetic segments isolated from an animal reservoir in Serbia. Archives of Biological Sciences, 2014, 66, 497-506.	0.2	1
39	Tracing the origin and dynamics of the HIV-1 epidemic in Serbia. Archives of Biological Sciences, 2014, 66, 507-515.	0.2	1
40	Neonatal herpes in Serbia: Is it a problem or not?. Archives of Biological Sciences, 2014, 66, 517-521.	0.2	0
41	Limited cross-border infections in patients newly diagnosed with HIV in Europe. Retrovirology, 2013, 10, 36.	0.9	52
42	HIV-1 subtype distribution and its demographic determinants in newly diagnosed patients in Europe suggest highly compartmentalized epidemics. Retrovirology, 2013, 10, 7.	0.9	129
43	International BioInformatics Workshop on Virus Evolution and Molecular Epidemiology. Infection, Genetics and Evolution, 2013, 19, 335-336.	1.0	1
44	Molecular typing of the local HIV-1 epidemic in Serbia. Infection, Genetics and Evolution, 2013, 19, 378-385.	1.0	15
45	mTOR-independent autophagy counteracts apoptosis in herpes simplex virus type 1-infected U251 glioma cells. Microbes and Infection, 2013, 15, 615-624.	1.0	30
46	p14ARF methylation is a common event in the pathogenesis and progression of myxoid and pleomorphic liposarcoma. Medical Oncology, 2013, 30, 682.	1.2	10
47	Treatment-associated polymorphisms in protease are significantly associated with higher viral load and lower CD4 count in newly diagnosed drug-naive HIV-1 infected patients. Retrovirology, 2012, 9, 81.	0.9	23
48	HIV-1 resistance profile in plasma and peripheral blood lymphocytes in a group of naive patients. Archives of Biological Sciences, 2012, 64, 1261-1270.	0.2	1
49	Predicting HIV treatment response in Romania – Comment. Germs, 2012, 2, 23-24.	O.5	2
50	HIV‑1 molecular epidemiology in the Balkans: a melting pot for high genetic diversity. AIDS Reviews, 2012, 14, 28-36.	0.5	29
51	Transmission of Drugâ€Resistant HIVâ€1 Is Stabilizing in Europe. Journal of Infectious Diseases, 2009, 200, 1503-1508.	1.9	213
52	Tracing the HIV-1 subtype B mobility in Europe: a phylogeographic approach. Retrovirology, 2009, 6, 49.	0.9	114
53	Functionally active virus-specific T cells that target CMV, adenovirus, and EBV can be expanded from naive T-cell populations in cord blood and will target a range of viral epitopes. Blood, 2009, 114, 1958-1967.	0.6	235
54	Disseminated Neonatal Herpes Caused by Herpes Simplex Virus Types 1 and 2. Emerging Infectious Diseases, 2007, 13, 302-304.	2.0	30

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#	Article	IF	CITATIONS
55	The Calculated Genetic Barrier for Antiretroviral Drug Resistance Substitutions Is Largely Similar for Different HIV-1 Subtypes. Journal of Acquired Immune Deficiency Syndromes (1999), 2006, 41, 352-360.	0.9	90
56	Synthesis, Characterization, and Biological Activity of Amino Acid Derivatives of the Heteropolytungstophosphoric Acid. Monatshefte Für Chemie, 2006, 137, 803-810.	0.9	11
57	Prevalence of Drugâ€Resistant HIVâ€1 Variants in Untreated Individuals in Europe: Implications for Clinical Management. Journal of Infectious Diseases, 2005, 192, 958-966.	1.9	385
58	Herpes simplex virus resistance to acyclovir in routine virological laboratory practice. Biomedicine and Pharmacotherapy, 2005, 59, 135-136.	2.5	1
59	HIV-1 Subtypes in Yugoslavia. AIDS Research and Human Retroviruses, 2002, 18, 519-522.	0.5	32
60	CMV DNA in blood and CSF of HIV infected patients. Virus Research, 2002, 85, 117-122.	1.1	3