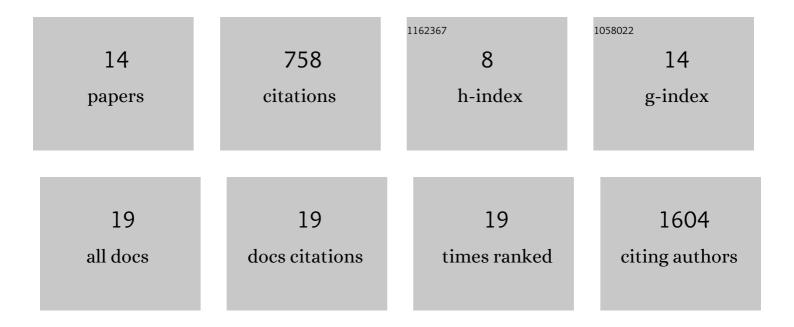
Anna Zhukova

List of Publications by Year in descending order

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ΔΝΝΑ ΖΗΠΚΟΛΑ

#	Article	IF	CITATIONS
1	Genome-wide diversity of Zika virus: Exploring spatio-temporal dynamics to guide a new nomenclature proposal. Virus Evolution, 2022, 8, veac029.	2.2	5
2	BioSimulators: a central registry of simulation engines and services for recommending specific tools. Nucleic Acids Research, 2022, 50, W108-W114.	6.5	11
3	Origin, evolution and global spread of SARS-CoV-2. Comptes Rendus - Biologies, 2021, 344, 57-75.	0.1	16
4	Maternal and neonatal outcomes related to Zika virus in pregnant women in Southern Vietnam: An epidemiological and virological prospective analysis. The Lancet Regional Health - Western Pacific, 2021, 11, 100163.	1.3	1
5	Phylogenetic analysis of HIV-1 shows frequent cross-country transmission and local population expansions. Virus Evolution, 2021, 7, veab055.	2.2	5
6	Cuban history of CRF19 recombinant subtype of HIV-1. PLoS Pathogens, 2021, 17, e1009786.	2.1	3
7	Drug resistance mutations in HIV: new bioinformatics approaches and challenges. Current Opinion in Virology, 2021, 51, 56-64.	2.6	23
8	The seventh pandemic of cholera in Europe revisited by microbial genomics. Nature Communications, 2020, 11, 5347.	5.8	13
9	<scp>SBML</scp> Level 3: an extensible format for the exchange and reuse of biological models. Molecular Systems Biology, 2020, 16, e9110.	3.2	178
10	A Fast Likelihood Method to Reconstruct and Visualize Ancestral Scenarios. Molecular Biology and Evolution, 2019, 36, 2069-2085.	3.5	153
11	The Role of Phylogenetics as a Tool to Predict the Spread of Resistance. Journal of Infectious Diseases, 2017, 216, S820-S823.	1.9	10
12	Toward Community Standards and Software for Whole-Cell Modeling. IEEE Transactions on Biomedical Engineering, 2016, 63, 2007-2014.	2.5	51
13	Pantograph: A template-based method for genome-scale metabolic model reconstruction. Journal of Bioinformatics and Computational Biology, 2015, 13, 1550006.	0.3	29
14	Controlled vocabularies and semantics in systems biology. Molecular Systems Biology, 2011, 7, 543.	3.2	246