

# Boris Vishnepolsky

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/368879/publications.pdf>

Version: 2024-02-01

10  
papers

714  
citations

1307594

7  
h-index

1474206

9  
g-index

11  
all docs

11  
docs citations

11  
times ranked

785  
citing authors

#	ARTICLE	IF	CITATIONS
1	DBAASP v3: database of antimicrobial/cytotoxic activity and structure of peptides as a resource for development of new therapeutics. <i>Nucleic Acids Research</i> , 2021, 49, D288-D297.	14.5	233
2	DBAASP v.2: an enhanced database of structure and antimicrobial/cytotoxic activity of natural and synthetic peptides. <i>Nucleic Acids Research</i> , 2016, 44, D1104-D1112.	14.5	169
3	<scp>dbaasp</scp>: database of antimicrobial activity and structure of peptides. <i>FEMS Microbiology Letters</i> , 2014, 357, 63-68.	1.8	83
4	Prediction of Linear Cationic Antimicrobial Peptides Based on Characteristics Responsible for Their Interaction with the Membranes. <i>Journal of Chemical Information and Modeling</i> , 2014, 54, 1512-1523.	5.4	67
5	Predictive Model of Linear Antimicrobial Peptides Active against Gram-Negative Bacteria. <i>Journal of Chemical Information and Modeling</i> , 2018, 58, 1141-1151.	5.4	57
6	Physicochemical Features and Peculiarities of Interaction of AMP with the Membrane. <i>Pharmaceuticals</i> , 2021, 14, 471.	3.8	46
7	De Novo Design and In Vitro Testing of Antimicrobial Peptides against Gram-Negative Bacteria. <i>Pharmaceuticals</i> , 2019, 12, 82.	3.8	42
8	Comparative analysis of machine learning algorithms on the microbial strain-specific AMP prediction. <i>Briefings in Bioinformatics</i> , 2022, 23, .	6.5	8
9	Comment on: "Empirical comparison of web-based antimicrobial peptide prediction tools". <i>Bioinformatics</i> , 2019, 35, 2692-2694.	4.1	7
10	Development of the model of in silico design of AMPs active against <i>Staphylococcus aureus</i> 25923. , 0, , .		2