

Khadijeh Ahmadi

List of Publications by Year in descending order

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Version: 2024-02-01

17
papers

295
citations

1684188

5
h-index

1125743

13
g-index

17
all docs

17
docs citations

17
times ranked

595
citing authors

#	ARTICLE	IF	CITATIONS
1	Characterization of SARS-CoV-2 isolated from a patient in Iran compared to SARS-CoV-2 different variants. <i>Journal of Biomolecular Structure and Dynamics</i> , 2023, 41, 1217-1232.	3.5	0
2	Enfuvirtide, an HIV-1 fusion inhibitor peptide, can act as a potent SARS-CoV-2 fusion inhibitor: an <i>in silico</i> drug repurposing study. <i>Journal of Biomolecular Structure and Dynamics</i> , 2022, 40, 5566-5576.	3.5	26
3	<i>In silico</i> identification of epitope-based vaccine candidates against HTLV-1. <i>Journal of Biomolecular Structure and Dynamics</i> , 2022, 40, 6737-6754.	3.5	7
4	Bioinformatics analyses for the designation of a hybrid protein against urinary tract infections caused by <i>Pseudomonas aeruginosa</i> and investigation of the presence of pre-existing antibodies in infected humans. <i>Journal of Biomolecular Structure and Dynamics</i> , 2022, 40, 9081-9095.	3.5	4
5	The Candidate Antigens to Achieving an Effective Vaccine against <i>Staphylococcus aureus</i> . <i>Vaccines</i> , 2022, 10, 199.	4.4	9
6	Vaccination of mice with hybrid protein containing Exotoxin S and PcrV with adjuvants alum and MPL protects <i>Pseudomonas aeruginosa</i> infections. <i>Scientific Reports</i> , 2022, 12, 1325.	3.3	2
7	Active site-based analysis of structural proteins for drug targets in different human <i>Coronaviruses</i> . <i>Chemical Biology and Drug Design</i> , 2022, 99, 585-602.	3.2	4
8	Reverse vaccinology approach to design a novel multi-epitope vaccine candidate against COVID-19: an <i>in silico</i> study. <i>Journal of Biomolecular Structure and Dynamics</i> , 2021, 39, 2857-2872.	3.5	187
9	Comparative study of the immune responses to the HMS-based fusion protein and capsule-based conjugated molecules as vaccine candidates in a mouse model of <i>Staphylococcus aureus</i> systemic infection. <i>Microbial Pathogenesis</i> , 2021, 150, 104656.	2.9	0
10	Protective efficacy of Hla-MntC-SACOL0723 fusion protein adjuvanted in alum and MPL against <i>Staphylococcus aureus</i> sepsis infection in mice. <i>Journal of Immunological Methods</i> , 2021, 494, 113055.	1.4	2
11	Evaluation of the immune response to a multi-epitope vaccine candidate in comparison with HlaH35L, MntC, and SACOL0723 proteins against MRSA infection. <i>Biologicals</i> , 2021, 73, 8-15.	1.4	2
12	<i>In silico</i> analysis of Single Nucleotide Polymorphisms Associated with MicroRNA Regulating 5-fluorouracil resistance in colorectal cancer. <i>Letters in Drug Design and Discovery</i> , 2021, 18, .	0.7	0
13	Virulence genes and antimicrobial resistance pattern in <i>Proteus mirabilis</i> strains isolated from patients attended with urinary infections to Tertiary Hospitals, in Iran. <i>African Health Sciences</i> , 2021, 21, 1677-84.	0.7	6
14	Preparation and preclinical evaluation of two novel <i>Staphylococcus aureus</i> capsular polysaccharide 5 and 8 fusion protein (Hla-MntC-SACOL0723) immunoconjugates. <i>IUBMB Life</i> , 2020, 72, 226-236.	3.4	5
15	<i>In silico</i> analysis and <i>in vivo</i> assessment of a novel epitope-based vaccine candidate against uropathogenic <i>Escherichia coli</i> . <i>Scientific Reports</i> , 2020, 10, 16258.	3.3	24
16	Epitope-based immunoinformatics study of a novel Hla-MntC-SACOL0723 fusion protein from <i>Staphylococcus aureus</i> : Induction of multi-pattern immune responses. <i>Molecular Immunology</i> , 2019, 114, 88-99.	2.2	12
17	Repurposing antiviral drugs against HTLV-1 protease by molecular docking and molecular dynamics simulation. <i>Journal of Biomolecular Structure and Dynamics</i> , 0, , 1-10.	3.5	5