

Akshay Uttarkar

List of Publications by Year in descending order

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Version: 2024-02-01

15
papers

199
citations

1162367

8
h-index

1125271

13
g-index

18
all docs

18
docs citations

18
times ranked

143
citing authors

#	ARTICLE	IF	CITATIONS
1	Structural and molecular basis of the interaction mechanism of selected drugs towards multiple targets of SARS-CoV-2 by molecular docking and dynamic simulation studies- deciphering the scope of repurposed drugs. <i>Computers in Biology and Medicine</i> , 2020, 126, 104054.	3.9	30
2	Carbon fullerene and nanotube are probable binders to multiple targets of SARS-CoV-2: Insights from computational modeling and molecular dynamic simulation studies. <i>Infection, Genetics and Evolution</i> , 2021, 96, 105155.	1.0	21
3	Structural insights on the interaction potential of natural leads against major protein targets of SARS-CoV-2: Molecular modelling, docking and dynamic simulation studies. <i>Computers in Biology and Medicine</i> , 2021, 132, 104325.	3.9	18
4	Coumarin derivative as a potent drug candidate against triple negative breast cancer targeting the frizzled receptor of wingless-related integration site signaling pathway. <i>Journal of Biomolecular Structure and Dynamics</i> , 2023, 41, 1561-1573.	2.0	18
5	Novel small molecules targeting bZIP23 TF improve stomatal conductance and photosynthesis under mild drought stress by regulating ABA. <i>FEBS Journal</i> , 2022, 289, 6058-6077.	2.2	17
6	Natural epiestriol-16 act as potential lead molecule against prospective molecular targets of multidrug resistant <i>Acinetobacter baumannii</i> -Insight from in silico modelling and in vitro investigations. <i>Infection, Genetics and Evolution</i> , 2020, 82, 104314.	1.0	16
7	Response regulator GacA and transcriptional activator RhIR proteins involved in biofilm formation of <i>Pseudomonas aeruginosa</i> are prospective targets for natural lead molecules: Computational modelling, molecular docking and dynamic simulation studies. <i>Infection, Genetics and Evolution</i> , 2020, 85, 104448.	1.0	13
8	Mitogen activated protein kinase-1 and cell division control protein-42 are putative targets for the binding of novel natural lead molecules: a therapeutic intervention against <i>Candida albicans</i> . <i>Journal of Biomolecular Structure and Dynamics</i> , 2020, 38, 4584-4599.	2.0	10
9	Spiro Benzodiazepine Substituted Fluorocoumarins as Potent Anti-Anxiety Agents. <i>Russian Journal of Bioorganic Chemistry</i> , 2021, 47, 390-398.	0.3	10
10	Stress-Induced Detoxification Enzymes in Rice Have Broad Substrate Affinity. <i>ACS Omega</i> , 2021, 6, 3399-3410.	1.6	9
11	Molecular Docking and Interaction Studies of Identified Abscisic Acid Receptors in <i>Oryza sativa</i> : An In-Silico Perspective on Comprehending Stress Tolerance Mechanisms. <i>Current Genomics</i> , 2021, 22, 607-619.	0.7	8
12	Understanding the Xylooligosaccharides Utilization Mechanism of <i>Lactobacillus brevis</i> and <i>Bifidobacterium adolescentis</i> : Proteins Involved and Their Conformational Stabilities for Effectual Binding. <i>Molecular Biotechnology</i> , 2022, 64, 75-89.	1.3	6
13	Carbon fullerene acts as potential lead molecule against prospective molecular targets of biofilm-producing multidrug-resistant <i>Acinetobacter baumannii</i> and <i>Pseudomonas aeruginosa</i> : computational modeling and MD simulation studies. <i>Journal of Biomolecular Structure and Dynamics</i> , 2021, 39, 1121-1137.	2.0	3
14	Study of SARS-nCoV2 Indian Isolates Gaining Insights into Mutation Frequencies, Protein Stability and Prospective Effect on Pathogenicity. <i>Coronaviruses</i> , 2021, 2, .	0.2	3
15	Genome Resequencing Reveals Single Nucleotide Polymorphism and Repeat Regions in <i>Giardia lamblia</i> Indian Isolate. <i>Journal of Next Generation Sequencing & Applications</i> , 2017, 04, .	0.3	1