

Dhurvas Chandrasekaran Dinesh

List of Publications by Year in descending order

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13
papers

851
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1597
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| # | ARTICLE | IF | CITATIONS |
|----|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----|-----------|
| 1 | Molecular Insights into Agonist/Antagonist Effects on Macromolecules Involved in Human Disease Mechanisms. <i>Current Molecular Pharmacology</i> , 2022, 15, 263-264. | 1.5 | 2 |
| 2 | High-Throughput Screening and Quantum Mechanics for Identifying Potent Inhibitors Against Mac1 Domain of SARS-CoV-2 Nsp3. <i>IEEE/ACM Transactions on Computational Biology and Bioinformatics</i> , 2021, 18, 1262-1270. | 3.0 | 17 |
| 3 | Structure-based virtual screening and molecular dynamics simulation of SARS-CoV-2 Guanine-N7 methyltransferase (nsp14) for identifying antiviral inhibitors against COVID-19. <i>Journal of Biomolecular Structure and Dynamics</i> , 2021, 39, 4582-4593. | 3.5 | 73 |
| 4 | Structural insights of macromolecules involved in bacteria-induced apoptosis in the pathogenesis of human diseases. <i>Advances in Protein Chemistry and Structural Biology</i> , 2021, 126, 1-38. | 2.3 | 5 |
| 5 | An <i>in silico</i> molecular dynamics simulation study on the inhibitors of SARS-CoV-2 proteases (3CL ^{pro} and PL ^{pro}) to combat COVID-19. <i>Molecular Simulation</i> , 2021, 47, 1168-1184. | 2.0 | 10 |
| 6 | Structural Understanding of SARS-CoV-2 Drug Targets, Active Site Contour Map Analysis and COVID-19 Therapeutics. <i>Current Molecular Pharmacology</i> , 2021, 14, . | 1.5 | 4 |
| 7 | Microsecond MD Simulation and Multiple-Conformation Virtual Screening to Identify Potential Anti-COVID-19 Inhibitors Against SARS-CoV-2 Main Protease. <i>Frontiers in Chemistry</i> , 2020, 8, 595273. | 3.6 | 32 |
| 8 | Structural basis of RNA recognition by the SARS-CoV-2 nucleocapsid phosphoprotein. <i>PLoS Pathogens</i> , 2020, 16, e1009100. | 4.7 | 206 |
| 9 | Antiviral Drug Targets of Single-Stranded RNA Viruses Causing Chronic Human Diseases. <i>Current Drug Targets</i> , 2020, 21, 105-124. | 2.1 | 18 |
| 10 | Structural Biology of Nuclear Auxin Action. <i>Trends in Plant Science</i> , 2016, 21, 302-316. | 8.8 | 45 |
| 11 | Solution structure of the PsIAA4 oligomerization domain reveals interaction modes for transcription factors in early auxin response. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 6230-6235. | 7.1 | 52 |
| 12 | Iron-Dependent Callose Deposition Adjusts Root Meristem Maintenance to Phosphate Availability. <i>Developmental Cell</i> , 2015, 33, 216-230. | 7.0 | 271 |
| 13 | Arabidopsis Calmodulin-binding Protein IQ67-Domain 1 Localizes to Microtubules and Interacts with Kinesin Light Chain-related Protein-1. <i>Journal of Biological Chemistry</i> , 2013, 288, 1871-1882. | 3.4 | 116 |