Surya Narayan Rath

List of Publications by Year in descending order

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1478505 1588992 10 107 6 8 citations g-index h-index papers 11 11 11 168 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Presentation of potential genes and deleterious variants associated with non-syndromic hearing loss: a computational approach. Genomics and Informatics, 2022, 20, e5.	0.8	O
2	Identification and evaluation of immunogenic MHC-I and MHC-II binding peptides from Mycobacterium tuberculosis. Computers in Biology and Medicine, 2021, 130, 104203.	7.0	6
3	In silico discovery and evaluation of phytochemicals binding mechanism against human catechol-O-methyltransferase as a putative bioenhancer of L-DOPA therapy in Parkinson disease. Genomics and Informatics, 2021, 19, e7.	0.8	3
4	Understanding ligands driven mechanism of wild and mutant aryl hydrocarbon receptor in presence of phytochemicals combating Parkinson's disease: an <i>in silico</i> and <i>in vivo</i> study. Journal of Biomolecular Structure and Dynamics, 2020, 38, 807-826.	3.5	8
5	Druggability for COVID-19: in silico discovery of potential drug compounds against nucleocapsid (N) protein of SARS-CoV-2. Genomics and Informatics, 2020, 18, e43.	0.8	13
6	<i>In silico</i> discovery of potential drug molecules to improve the treatment of isoniazid-resistant <i>Mycobacterium tuberculosis</i> Dynamics, 2019, 37, 3388-3398.	3.5	14
7	Identification of Suitable Natural Inhibitor against Influenza A (H1N1) Neuraminidase Protein by Molecular Docking. Genomics and Informatics, 2016, 14, 96.	0.8	39
8	Drug Target Identification and Elucidation of Natural Inhibitors for <i>Bordetella petrii </i> Silico Study. Genomics and Informatics, 2016, 14, 241.	0.8	10
9	Elucidation of the Inhibitory Effect of Phytochemicals with Kir6.2 Wild-Type and Mutant Models Associated in Type-1 Diabetes through Molecular Docking Approach. Genomics and Informatics, 2014, 12, 283.	0.8	10
10	Structural Analysis of Respirasomes in Electron Transfer Pathway of <i>Acidithiobacillus ferrooxidans</i> : A Computer-Aided Molecular Designing Study., 2013, 2013, 1-14.		4