## Dhanusha Yesudhas

List of Publications by Year in descending order

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933264 887953 18 476 10 17 citations g-index h-index papers 18 18 18 816 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	COVID-19 outbreak: history, mechanism, transmission, structural studies and therapeutics. Infection, 2021, 49, 199-213.	2.3	160
2	Toll-like receptors: promising therapeutic targets for inflammatory diseases. Archives of Pharmacal Research, 2016, 39, 1032-1049.	2.7	80
3	Multiple Roles of Toll-Like Receptor 4 in Colorectal Cancer. Frontiers in Immunology, 2014, 5, 334.	2.2	77
4	Proteins Recognizing DNA: Structural Uniqueness and Versatility of DNA-Binding Domains in Stem Cell Transcription Factors. Genes, 2017, 8, 192.	1.0	28
5	A Combined Molecular Docking/Dynamics Approach to Probe the Binding Mode of Cancer Drugs with Cytochrome P450 3A4. Molecules, 2015, 20, 14915-14935.	1.7	22
6	Structural Mechanism behind Distinct Efficiency of Oct4/Sox2 Proteins in Differentially Spaced DNA Complexes. PLoS ONE, 2016, 11, e0147240.	1.1	19
7	Evaluation of Sox2 binding affinities for distinct <scp>DNA</scp> patterns using steered molecular dynamics simulation. FEBS Open Bio, 2017, 7, 1750-1767.	1.0	18
8	Tollâ€like receptor 2 antagonists identified through virtual screening and experimental validation. FEBS Journal, 2017, 284, 2264-2283.	2.2	17
9	VEPAD - Predicting the effect of variants associated with Alzheimer's disease using machine learning. Computers in Biology and Medicine, 2020, 124, 103933.	3.9	14
10	PAMAM/5-fluorouracil drug conjugate for targeting E6 and E7 oncoproteins in cervical cancer: a combined experimental/in silico approach. RSC Advances, 2017, 7, 5046-5054.	1.7	13
11	Tackling Covidâ€19 using disorderedâ€toâ€order transition of residues in the spike protein upon angiotensinâ€converting enzyme 2 binding. Proteins: Structure, Function and Bioinformatics, 2021, 89, 1158-1166.	1.5	9
12	Cysteine redox state plays a key role in the inter-domain movements of HMGB1: a molecular dynamics simulation study. RSC Advances, 2016, 6, 100804-100819.	1.7	6
13	Structural and conformational insights into SOX2/OCT4-bound enhancer DNA: a computational perspective. RSC Advances, 2016, 6, 90138-90153.	1.7	4
14	Role of disordered regions in transferring tyrosine to its cognate tRNA. International Journal of Biological Macromolecules, 2020, 150, 705-713.	3.6	3
15	Understanding disorder-to-order transitions in protein–RNA complexes using molecular dynamics simulations. Journal of Biomolecular Structure and Dynamics, 2021, , 1-11.	2.0	3
16	Tumor Heterogeneity and Molecular Characteristics of Glioblastoma Revealed by Single-Cell RNA-Seq Data Analysis. Genes, 2022, 13, 428.	1.0	2
17	Structural mechanism of DNA-mediated Nanog–Sox2 cooperative interaction. RSC Advances, 2019, 9, 8121-8130.	1.7	1
18	Deciphering the Role of Residues Involved in Disorder-To-Order Transition Regions in Archaeal tRNA Methyltransferase 5. Genes, 2021, 12, 399.	1.0	0