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List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2233197/publications.pdf>

Version: 2024-02-01

12
papers

329
citations

1040056

9
h-index

1281871

11
g-index

12
all docs

12
docs citations

12
times ranked

514
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic variants of small airways and interstitial pulmonary disease in children. <i>Scientific Reports</i> , 2021, 11, 2715.	3.3	4
2	Role of SARS-CoV-2 and ACE2 variations in COVID-19. <i>Biomedical Journal</i> , 2021, 44, 235-244.	3.1	20
3	Molecular Dynamics Simulation Study of the Interaction between Human Angiotensin Converting Enzyme 2 and Spike Protein Receptor Binding Domain of the SARS-CoV-2 B.1.617 Variant. <i>Biomolecules</i> , 2021, 11, 1244.	4.0	17
4	Bioactive Peptides as Potential Nutraceuticals for Diabetes Therapy: A Comprehensive Review. <i>International Journal of Molecular Sciences</i> , 2021, 22, 9059.	4.1	35
5	Genetic variants in children with chronic respiratory diseases. <i>Pediatric Pulmonology</i> , 2020, 55, 2389-2401.	2.0	5
6	Antioxidant and anticancer properties of berries. <i>Critical Reviews in Food Science and Nutrition</i> , 2018, 58, 2491-2507.	10.3	130
7	Interactions of quercetin with receptor tyrosine kinases associated with human lung carcinoma. <i>Natural Product Research</i> , 2018, 32, 2928-2931.	1.8	20
8	Structural insights into the polypharmacological activity of quercetin on serine/threonine kinases. <i>Drug Design, Development and Therapy</i> , 2016, Volume 10, 3109-3123.	4.3	19
9	Acetogenins from <i>Annona muricata</i> as potential inhibitors of antiapoptotic proteins: a molecular modeling study. <i>Drug Design, Development and Therapy</i> , 2016, 10, 1399.	4.3	14
10	Molecular insights into the binding of phosphoinositides to the TH domain region of TIPE proteins. <i>Journal of Molecular Modeling</i> , 2016, 22, 272.	1.8	15
11	Identification of Novel Aldose Reductase Inhibitors from Spices: A Molecular Docking and Simulation Study. <i>PLoS ONE</i> , 2015, 10, e0138186.	2.5	49
12	Dynamics of the interaction between the receptor-binding domain of SARS-CoV-2 Omicron (B.1.1.529) variant and human angiotensin-converting enzyme 2. <i>PeerJ</i> , 0, 10, e13680.	2.0	1