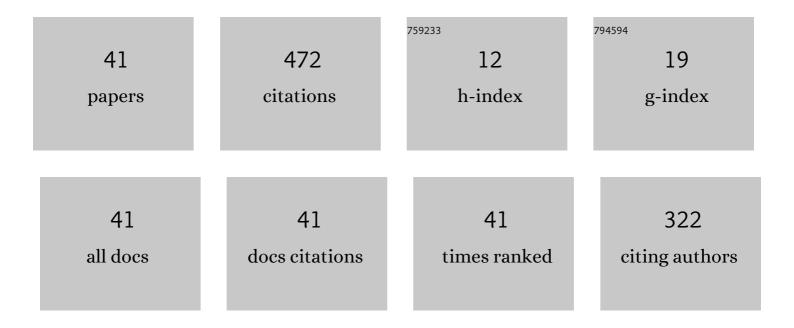
Sani Uba

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Molecular Docking , ADMET and Pharmacokinetic properties predictions of some di-aryl Pyridinamine derivatives as Estrogen Receptor (Er+) Kinase Inhibitors. Egyptian Journal of Basic and Applied Sciences, 2022, 9, 180-204.	0.6	4
2	QSAR, Ligand Based Design and Pharmacokinetic Studies of Parviflorons Derivatives as Anti-Breast Cancer Drug Compounds Against MCF-7 Cell Line. Chemistry Africa, 2021, 4, 175-187.	2.4	4
3	Computer modeling of some anti-breast cancer compounds. Structural Chemistry, 2021, 32, 679-687.	2.0	6
4	Ligand-based drug design and molecular docking simulation studies of some novel anticancer compounds on MALME-3M melanoma cell line. Egyptian Journal of Medical Human Genetics, 2021, 22, .	1.0	8
5	MOLECULAR DOCKING INVESTIGATION AND PHARMACOKINETIC PROPERTIES PREDICTION OF SOME ANILINOPYRIMIDINES ANALOGUES AS EGFR T790M TYROSINE KINASE INHIBITORS. Egyptian Journal of Basic and Applied Sciences, 2021, 8, 203-213.	0.6	3
6	Conceptual Structure-Based Drug Design and Discovering of Novel Inhibitors of Norepinephrine Transporter (NET) as Potential Antipsychotic Agents for Mental Disorder. Chemistry Africa, 2021, 4, 115-125.	2.4	0
7	Unveiling novel inhibitors of dopamine transporter via in silico drug design, molecular docking, and bioavailability predictions as potential antischizophrenic agents. Future Journal of Pharmaceutical Sciences, 2021, 7, .	2.8	10
8	Computer-aided design of some quinazoline analogues as epidermal growth factor receptor inhibitors. Egyptian Journal of Medical Human Genetics, 2021, 22, .	1.0	1
9	Design of more potent quinazoline derivatives as EGFRWT inhibitors for the treatment of NSCLC: a computational approach. Future Journal of Pharmaceutical Sciences, 2021, 7, .	2.8	9
10	Computer-aided drug design and in silico pharmacokinetics predictions of some potential antipsychotic agents. Scientific African, 2021, 12, e00734.	1.5	5
11	In silico study for evaluating the binding mode and interaction of 1, 2, 4-triazole and its derivatives as potent inhibitors against Lipoate protein B (LipB). Journal of King Saud University - Science, 2020, 32, 475-485.	3.5	11
12	Multi-linear regression model, molecular binding interactions and ligand-based design of some prominent compounds against Mycobacterium tuberculosis. Network Modeling Analysis in Health Informatics and Bioinformatics, 2020, 9, 1.	2.1	8
13	Structure-based design and activity modeling of novel epidermal growth factor receptor kinase inhibitors; an in silico approach. Scientific African, 2020, 9, e00503.	1.5	13
14	In Silico Modeling, Prediction, and Designing of Some Anti-wear Lubricant Additives. Journal of Bio- and Tribo-Corrosion, 2020, 6, 1.	2.6	1
15	QSAR, molecular docking, design, and pharmacokinetic analysis of 2-(4-fluorophenyl) imidazol-5-ones as anti-breast cancer drug compounds against MCF-7 cell line. Journal of Bioenergetics and Biomembranes, 2020, 52, 475-494.	2.3	3
16	Chemoinformatic studies on some inhibitors of dopamine transporter and the receptor targeting schizophrenia for developing novel antipsychotic agents. Heliyon, 2020, 6, e04464.	3.2	8
17	Lead Identification of Some Anti-Cancer Agents with Prominent Activity Against Non-small Cell Lung Cancer (NSCLC) and Structure-Based Design. Chemistry Africa, 2020, 3, 1023-1044.	2.4	13
18	Profiling the antidepressant properties of phenyl piperidine derivatives as inhibitors of serotonin transporter (SERT) via cheminformatics modeling, molecular docking and ADMET predictions Scientific African, 2020, 9, e00517.	1.5	5

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19	Design of potential anti-melanoma agents against SK-MEL-5 cell line using QSAR modeling and molecular docking methods. SN Applied Sciences, 2020, 2, 1.	2.9	22
20	Computational Modeling and Pharmacokinetics/ADMET Study of Some Arylpiperazine Derivatives as Novel Antipsychotic Agents Targeting Depression. Chemistry Africa, 2020, 3, 979-988.	2.4	6
21	Computational modeling of novel quinazoline derivatives as potent epidermal growth factor receptor inhibitors. Heliyon, 2020, 6, e03289.	3.2	18
22	In-silico activity prediction and docking studies of some 2, 9-disubstituted 8-phenylthio/phenylsulfinyl-9h-purine derivatives as Anti-proliferative agents. Heliyon, 2020, 6, e03158.	3.2	34
23	QSAR modeling, molecular docking and ADMET/pharmacokinetic studies: a chemometrics approach to search for novel inhibitors of norepinephrine transporter as potent antipsychotic drugs. Journal of the Iranian Chemical Society, 2020, 17, 1953-1966.	2.2	19
24	Computational studies of a series of 2-substituted phenyl-2-oxo-, 2-hydroxyl- and 2-acylloxyethylsulfonamides as potent anti-fungal agents. Heliyon, 2020, 6, e03724.	3.2	49
25	Structure-based design of some quinazoline derivatives as epidermal growth factor receptor inhibitors. Egyptian Journal of Medical Human Genetics, 2020, 21, .	1.0	4
26	In silico studies of some 2-anilinopyrimidine derivatives as anti-triple-negative breast cancer agents. Beni-Suef University Journal of Basic and Applied Sciences, 2020, 9, .	2.0	3
27	Computational virtual screening and structure-based design of some epidermal growth factor receptor inhibitors. Future Journal of Pharmaceutical Sciences, 2020, 6, .	2.8	5
28	Quantitative structure-activity relationship, molecular docking, drug-likeness, and pharmacokinetic studies of some non-small cell lung cancer therapeutic agents. Beni-Suef University Journal of Basic and Applied Sciences, 2020, 9, .	2.0	6
29	QSAR analysis and molecular docking simulation of norepinephrine transporter (NET) inhibitors as anti-psychotic therapeutic agents. Heliyon, 2019, 5, e02640.	3.2	16
30	A Derived QSAR Model for Predicting Some Compounds as Potent Antagonist against <i> Mycobacterium tuberculosis</i> : A Theoretical Approach. Advances in Preventive Medicine, 2019, 2019, 1-18.	2.7	24
31	Molecular design and docking analysis of the inhibitory activities of some α_substituted acetamido-N-benzylacetamide as anticonvulsant agents. SN Applied Sciences, 2019, 1, 1.	2.9	9
32	Molecular design of antioxidant lubricating oil additives via QSPR and analysis dynamic simulation method. Heliyon, 2019, 5, e02880.	3.2	7
33	QSAR modelling and docking analysis of some thiazole analogues as alfa-glucosidase inhibitors. The Journal of Engineering and Exact Sciences, 2019, 5, 0257-0270.	0.1	8
34	QSAR STUDY OF 2-SUBSTITUTED PHENYL-2-OXO-, 2-HYDROXYL- AND 2-ACYLLOXYETHYLSULFONAMIDES AS FUNGICIDES. The Journal of Engineering and Exact Sciences, 2019, 5, 0283-0290.	0.1	3
35	Molecular docking and quantitative structure-activity relationship study of anticonvulsant activity of aminobenzothiazole derivatives. Beni-Suef University Journal of Basic and Applied Sciences, 2018, 7, 204-214.	2.0	13
36	Theoretical modeling and molecular docking simulation for investigating and evaluating some active compounds as potent anti-tubercular agents against MTB CYP121 receptor. Future Journal of Pharmaceutical Sciences, 2018, 4, 284-295.	2.8	24

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37	QSAR Modeling and Molecular Docking Analysis of Some Active Compounds against <i> Mycobacterium tuberculosis</i> Receptor (Mtb CYP121). Journal of Pathogens, 2018, 2018, 1-24.	1.4	28
38	IN SILICO STUDY FOR INVESTIGATING AND PREDICTING THE ACTIVITIES OF 1,2,4-TRIAZOLE DERIVATIES AS POTENT ANTI-TUBERCULAR AGENTS. The Journal of Engineering and Exact Sciences, 2018, 4, 0246-0254.	0.1	7
39	Quantitative structure-activity relationship and molecular docking studies of a series of quinazolinonyl analogues as inhibitors of gamma amino butyric acid aminotransferase. Journal of Advanced Research, 2017, 8, 33-43.	9.5	37
40	Quantitative structure activity relationship study of anticonvulsant activity of $\hat{1}\pm$ _substituted acetamido-N-benzylacetamide derivatives. Cogent Chemistry, 2016, 2, 1166538.	2.5	11
41	Effect of Acetylation on Stability to Retrogradation of Starch Extracted from Wild Polynesian Arrowroot (<i>Tacca leontopetaloides</i> (L.) Kuntze) for Utilization as Adhesive on Paper. Journal of Polymers, 2014, 2014, 1-9.	0.9	7