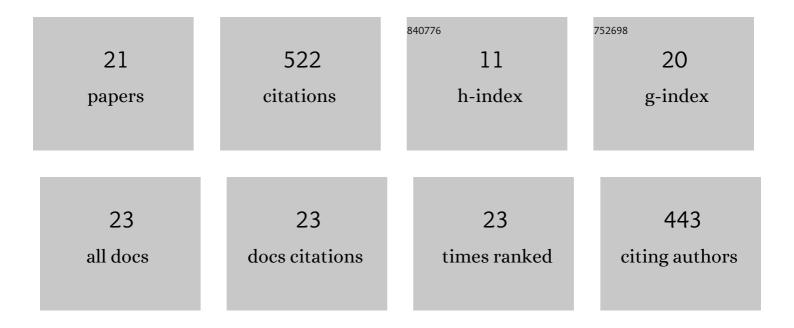
## Abdus Samad

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9257810/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Designing a multi-epitope vaccine against SARS-CoV-2: an immunoinformatics approach. Journal of Biomolecular Structure and Dynamics, 2022, 40, 14-30.	3.5	113
2	A systematic analysis of ATPase Cation transporting 13A2 (ATP13A2) transcriptional expression and prognostic value in human brain cancer. Biomedical Signal Processing and Control, 2022, 71, 103183.	5.7	3
3	Evaluation of in vitro and in silico anti-inflammatory potential of some selected medicinal plants of Bangladesh against cyclooxygenase-II enzyme. Journal of Ethnopharmacology, 2022, 285, 114900.	4.1	19
4	Bioinformatics approaches identified dasatinib and bortezomib inhibit the activity of MCM7 protein as a potential treatment against human cancer. Scientific Reports, 2022, 12, 1539.	3.3	16
5	Computational Identification of Druggable Bioactive Compounds from Catharanthus roseus and Avicennia marina against Colorectal Cancer by Targeting Thymidylate Synthase. Molecules, 2022, 27, 2089.	3.8	19
6	Immune epitopes identification and designing of a multi-epitope vaccine against bovine leukemia virus: a molecular dynamics and immune simulation approaches. Cancer Immunology, Immunotherapy, 2022, 71, 2535-2548.	4.2	12
7	Toward the Identification of Natural Antiviral Drug Candidates against Merkel Cell Polyomavirus: Computational Drug Design Approaches. Pharmaceuticals, 2022, 15, 501.	3.8	7
8	High expression of bone morphogenetic protein 1 (BMP1) is associated with a poor survival rate in human gastric cancer, a dataset approaches. Genomics, 2021, 113, 1141-1154.	2.9	19
9	Methanol, ethyl acetate and n-hexane extracts of Tragia involucrata L. leaves exhibit anxiolytic, sedative and analgesic activity in Swiss albino mice. Heliyon, 2021, 7, e05814.	3.2	4
10	Pharmacoinformatics and molecular dynamics simulation-based phytochemical screening of neem plant <i>(Azadiractha indica)</i> against human cancer by targeting MCM7 protein. Briefings in Bioinformatics, 2021, 22, .	6.5	55
11	Transporter associated with antigen processing 1 (TAP1) expression and prognostic analysis in breast, lung, liver, and ovarian cancer. Journal of Molecular Medicine, 2021, 99, 1293-1309.	3.9	29
12	Transcriptomic studies revealed pathophysiological impact of COVID-19 to predominant health conditions. Briefings in Bioinformatics, 2021, 22, .	6.5	17
13	Pharmacophore-Based Virtual Screening, Quantum Mechanics Calculations, and Molecular Dynamics Simulation Approaches Identified Potential Natural Antiviral Drug Candidates against MERS-CoV S1-NTD. Molecules, 2021, 26, 4961.	3.8	36
14	Spike protein recognizer receptor ACE2 targeted identification of potential natural antiviral drug candidates against SARS-CoV-2. International Journal of Biological Macromolecules, 2021, 191, 1114-1125.	7.5	36
15	Identification of angiotensin-converting enzyme 2 (ACE2) protein as the potential biomarker in SARS-CoV-2 infection-related lung cancer using computational analyses. Genomics, 2020, 112, 4912-4923.	2.9	31
16	A multi-omics approach to reveal the key evidence of GDF10 as a novel therapeutic biomarker for breast cancer. Informatics in Medicine Unlocked, 2020, 21, 100463.	3.4	9
17	Clinicopathological and prognostic significance of GPC3 in human breast cancer and its 3D structure prediction. Network Modeling Analysis in Health Informatics and Bioinformatics, 2020, 9, 1.	2.1	9
18	A Multi-Omics Analysis of Bone Morphogenetic Protein 5 (BMP5) mRNA Expression and Clinical Prognostic Outcomes in Different Cancers Using Bioinformatics Approaches. Biomedicines, 2020, 8, 19.	3.2	23

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#	Article	IF	CITATIONS
19	Validation of CSN1S1 transcriptional expression, promoter methylation, and prognostic power in breast cancer using independent datasets. Biochemistry and Biophysics Reports, 2020, 24, 100867.	1.3	10
20	Computational assessment of MCM2 transcriptional expression and identification of the prognostic biomarker for human breast cancer. Heliyon, 2020, 6, e05087.	3.2	34
21	Discovery of potential immune epitopes and peptide vaccine design - a prophylactic strategy against Rift Valley fever virus. F1000Research, 0, 9, 999.	1.6	17