

# Taj Mohammad

## List of Publications by Year in descending order

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Version: 2024-02-01

81  
papers

3,501  
citations

147566

31  
h-index

161609

54  
g-index

82  
all docs

82  
docs citations

82  
times ranked

3029  
citing authors

#	ARTICLE	IF	CITATIONS
1	Insights into SARS-CoV-2 genome, structure, evolution, pathogenesis and therapies: Structural genomics approach. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2020, 1866, 165878.	1.8	770
2	Advancements in Docking and Molecular Dynamics Simulations Towards Ligand-receptor Interactions and Structure-function Relationships. <i>Current Topics in Medicinal Chemistry</i> , 2018, 18, 1755-1768.	1.0	188
3	Glecaprevir and Maraviroc are high-affinity inhibitors of SARS-CoV-2 main protease: possible implication in COVID-19 therapy. <i>Bioscience Reports</i> , 2020, 40, .	1.1	129
4	InstaDock: A single-click graphical user interface for molecular docking-based virtual high-throughput screening. <i>Briefings in Bioinformatics</i> , 2021, 22, .	3.2	127
5	Identification and evaluation of bioactive natural products as potential inhibitors of human microtubule affinity-regulating kinase 4 (MARK4). <i>Journal of Biomolecular Structure and Dynamics</i> , 2019, 37, 1813-1829.	2.0	114
6	Investigation of molecular mechanism of recognition between citral and MARK4: A newer therapeutic approach to attenuate cancer cell progression. <i>International Journal of Biological Macromolecules</i> , 2018, 107, 2580-2589.	3.6	96
7	Virtual Screening Approach to Identify High-Affinity Inhibitors of Serum and Glucocorticoid-Regulated Kinase 1 among Bioactive Natural Products: Combined Molecular Docking and Simulation Studies. <i>Molecules</i> , 2020, 25, 823.	1.7	89
8	Investigation of inhibitory potential of quercetin to the pyruvate dehydrogenase kinase 3: Towards implications in anticancer therapy. <i>International Journal of Biological Macromolecules</i> , 2019, 136, 1076-1085.	3.6	80
9	Identification of high-affinity inhibitors of SARS-CoV-2 main protease: Towards the development of effective COVID-19 therapy. <i>Virus Research</i> , 2020, 288, 198102.	1.1	79
10	Evaluation of ellagic acid as an inhibitor of sphingosine kinase 1: A targeted approach towards anticancer therapy. <i>Biomedicine and Pharmacotherapy</i> , 2019, 118, 109245.	2.5	78
11	Potential drug targets of SARS-CoV-2: From genomics to therapeutics. <i>International Journal of Biological Macromolecules</i> , 2021, 177, 1-9.	3.6	77
12	Discovery of Hordenine as a Potential Inhibitor of Pyruvate Dehydrogenase Kinase 3: Implication in Lung Cancer Therapy. <i>Biomedicines</i> , 2020, 8, 119.	1.4	76
13	MARK4 Inhibited by AChE Inhibitors, Donepezil and Rivastigmine Tartrate: Insights into Alzheimer's Disease Therapy. <i>Biomolecules</i> , 2020, 10, 789.	1.8	76
14	Targeting pyruvate dehydrogenase kinase signaling in the development of effective cancer therapy. <i>Biochimica Et Biophysica Acta: Reviews on Cancer</i> , 2021, 1876, 188568.	3.3	75
15	Identification and Evaluation of Inhibitors of Lipase from <i>Malassezia restricta</i> using Virtual High-Throughput Screening and Molecular Dynamics Studies. <i>International Journal of Molecular Sciences</i> , 2019, 20, 884.	1.8	72
16	Identification of $\beta$ -Mangostin as a Potential Inhibitor of Microtubule Affinity Regulating Kinase 4. <i>Journal of Natural Products</i> , 2019, 82, 2252-2261.	1.5	62
17	FNDC5/Irisin: Physiology and Pathophysiology. <i>Molecules</i> , 2022, 27, 1118.	1.7	56
18	Structural Analysis and Conformational Dynamics of STN1 Gene Mutations Involved in Coat Plus Syndrome. <i>Frontiers in Molecular Biosciences</i> , 2019, 6, 41.	1.6	53

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19	Evaluation of binding and inhibition mechanism of dietary phytochemicals with sphingosine kinase 1: Towards targeted anticancer therapy. <i>Scientific Reports</i> , 2019, 9, 18727.	1.6	52
20	Probing the interaction of Rivastigmine Tartrate, an important Alzheimer's drug, with serum albumin: Attempting treatment of Alzheimer's disease. <i>International Journal of Biological Macromolecules</i> , 2020, 148, 533-542.	3.6	52
21	Identification of natural compounds as potent inhibitors of SARS-CoV-2 main protease using combined docking and molecular dynamics simulations. <i>Saudi Journal of Biological Sciences</i> , 2021, 28, 2423-2431.	1.8	47
22	Unraveling Binding Mechanism of Alzheimer's Drug Rivastigmine Tartrate with Human Transferrin: Molecular Docking and Multi-Spectroscopic Approach towards Neurodegenerative Diseases. <i>Biomolecules</i> , 2019, 9, 495.	1.8	46
23	Mutated CEACAMs Disrupt Transforming Growth Factor Beta Signaling and Alter the Intestinal Microbiome to Promote Colorectal Carcinogenesis. <i>Gastroenterology</i> , 2020, 158, 238-252.	0.6	46
24	Identification and evaluation of glutathione conjugate gamma-glutamyl-cysteine for improved drug delivery to the brain. <i>Journal of Biomolecular Structure and Dynamics</i> , 2020, 38, 3610-3620.	2.0	46
25	Exploring molecular insights into the interaction mechanism of cholesterol derivatives with the Mce4A: A combined spectroscopic and molecular dynamic simulation studies. <i>International Journal of Biological Macromolecules</i> , 2018, 111, 548-560.	3.6	45
26	Molecular interaction studies on ellagic acid for its anticancer potential targeting pyruvate dehydrogenase kinase 3. <i>RSC Advances</i> , 2019, 9, 23302-23315.	1.7	44
27	Genomic Variations in the Structural Proteins of SARS-CoV-2 and Their Deleterious Impact on Pathogenesis: A Comparative Genomics Approach. <i>Frontiers in Cellular and Infection Microbiology</i> , 2021, 11, 765039.	1.8	43
28	B Cell Lymphoma 2: A Potential Therapeutic Target for Cancer Therapy. <i>International Journal of Molecular Sciences</i> , 2021, 22, 10442.	1.8	39
29	Identification of Sphingosine Kinase-1 Inhibitors from Bioactive Natural Products Targeting Cancer Therapy. <i>ACS Omega</i> , 2020, 5, 14720-14729.	1.6	38
30	Vitamin D and lumisterol novel metabolites can inhibit SARS-CoV-2 replication machinery enzymes. <i>American Journal of Physiology - Endocrinology and Metabolism</i> , 2021, 321, E246-E251.	1.8	38
31	Identification of High-Affinity Inhibitors of Cyclin-Dependent Kinase 2 Towards Anticancer Therapy. <i>Molecules</i> , 2019, 24, 4589.	1.7	37
32	Discovery of Harmaline as a Potent Inhibitor of Sphingosine Kinase-1: A Chemopreventive Role in Lung Cancer. <i>ACS Omega</i> , 2020, 5, 21550-21560.	1.6	35
33	Investigation of deleterious effects of nsSNPs in the POT1 gene: a structural genomics-based approach to understand the mechanism of cancer development. <i>Journal of Cellular Biochemistry</i> , 2019, 120, 10281-10294.	1.2	32
34	Virtual high-throughput screening of natural compounds in-search of potential inhibitors for protection of telomeres 1 (POT1). <i>Journal of Biomolecular Structure and Dynamics</i> , 2020, 38, 4625-4634.	2.0	32
35	Phytoconstituents and Medicinal Plants for Anticancer Drug Discovery: Computational Identification of Potent Inhibitors of PIM1 Kinase. <i>OMICS A Journal of Integrative Biology</i> , 2021, 25, 580-590.	1.0	30
36	Impact of Gln94Glu mutation on the structure and function of protection of telomere 1, a cause of cutaneous familial melanoma. <i>Journal of Biomolecular Structure and Dynamics</i> , 2020, 38, 1514-1524.	2.0	29

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37	Evaluation of Binding of Rosmarinic Acid with Human Transferrin and Its Impact on the Protein Structure: Targeting Polyphenolic Acid-Induced Protection of Neurodegenerative Disorders. <i>Oxidative Medicine and Cellular Longevity</i> , 2020, 2020, 1-14.	1.9	26
38	Identification of Phytoconstituents as Potent Inhibitors of Casein Kinase-1 Alpha Using Virtual Screening and Molecular Dynamics Simulations. <i>Pharmaceutics</i> , 2021, 13, 2157.	2.0	25
39	Identification of plant-based hexokinase 2 inhibitors: combined molecular docking and dynamics simulation studies. <i>Journal of Biomolecular Structure and Dynamics</i> , 2022, 40, 10319-10331.	2.0	23
40	Î2-spectrin (SPTBN1) as a therapeutic target for diet-induced liver disease and preventing cancer development. <i>Science Translational Medicine</i> , 2021, 13, eabk2267.	5.8	23
41	Identification of high-affinity inhibitors of pyruvate dehydrogenase kinase-3: towards therapeutic management of cancer. <i>Journal of Biomolecular Structure and Dynamics</i> , 2021, 39, 586-594.	2.0	22
42	Thymoquinone and quercetin induce enhanced apoptosis in non-small cell lung cancer in combination through the Bax/Bcl2 cascade. <i>Journal of Cellular Biochemistry</i> , 2022, 123, 259-274.	1.2	20
43	Insight into the binding of PEG-400 with eye protein alpha-crystallin: Multi spectroscopic and computational approach: possible therapeutics targeting eye diseases. <i>Journal of Biomolecular Structure and Dynamics</i> , 2022, 40, 4496-4506.	2.0	19
44	Impact of amino acid substitution in the kinase domain of Bruton tyrosine kinase and its association with X-linked agammaglobulinemia. <i>International Journal of Biological Macromolecules</i> , 2020, 164, 2399-2408.	3.6	18
45	Investigation of conformational dynamics of Tyr89Cys mutation in protection of telomeres 1 gene associated with familial melanoma. <i>Journal of Biomolecular Structure and Dynamics</i> , 2021, 39, 35-44.	2.0	18
46	Structural genomics approach to investigate deleterious impact of nsSNPs in conserved telomere maintenance component 1. <i>Scientific Reports</i> , 2021, 11, 10202.	1.6	18
47	Structural and functional impact of non-synonymous SNPs in the CST complex subunit TEN1: structural genomics approach. <i>Bioscience Reports</i> , 2019, 39, .	1.1	17
48	Identification and structure-activity relationship (SAR) studies of carvacrol derivatives as potential anti-malarial against Plasmodium falciparum falcipain-2 protease. <i>Bioorganic Chemistry</i> , 2020, 103, 104142.	2.0	16
49	Impact of non-synonymous mutations on the structure and function of telomeric repeat binding factor 1. <i>Journal of Biomolecular Structure and Dynamics</i> , 2022, 40, 9053-9066.	2.0	16
50	Insights into the Conserved Regulatory Mechanisms of Human and Yeast Aging. <i>Biomolecules</i> , 2020, 10, 882.	1.8	15
51	Bioactive phytoconstituents as potent inhibitors of casein kinase-2: dual implications in cancer and COVID-19 therapeutics. <i>RSC Advances</i> , 2022, 12, 7872-7882.	1.7	14
52	Design and Development of Novel Urea, Sulfonylurea, and Sulfonamide Derivatives as Potential Inhibitors of Sphingosine Kinase 1. <i>Pharmaceutics</i> , 2020, 13, 118.	1.7	13
53	Multispectroscopic and Molecular Docking Insight into Elucidating the Interaction of Irisin with Rivastigmine Tartrate: A Combinational Therapy Approach to Fight Alzheimer's Disease. <i>ACS Omega</i> , 2021, 6, 7910-7921.	1.6	13
54	Investigating single amino acid substitutions in PIM1 kinase: A structural genomics approach. <i>PLoS ONE</i> , 2021, 16, e0258929.	1.1	13

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55	Aurora B kinase: a potential drug target for cancer therapy. <i>Journal of Cancer Research and Clinical Oncology</i> , 2021, 147, 2187-2198.	1.2	12
56	Discovery of Natural Compounds as Potential Inhibitors of Human Carbonic Anhydrase II: An Integrated Virtual Screening, Docking, and Molecular Dynamics Simulation Study. <i>OMICS A Journal of Integrative Biology</i> , 2021, 25, 513-524.	1.0	12
57	Impact of Deleterious Mutations on Structure, Function and Stability of Serum/Glucocorticoid Regulated Kinase 1: A Gene to Diseases Correlation. <i>Frontiers in Molecular Biosciences</i> , 2021, 8, 780284.	1.6	12
58	Impact of single amino acid substitution on the structure and function of TANK-binding kinase-1. <i>Journal of Cellular Biochemistry</i> , 2021, 122, 1475-1490.	1.2	11
59	Identification of intrinsically disorder regions in non-structural proteins of SARS-CoV-2: New insights into drug and vaccine resistance. <i>Molecular and Cellular Biochemistry</i> , 2022, 477, 1607-1619.	1.4	11
60	Inhibiting Cyclin-Dependent Kinase 6 by Taurine: Implications in Anticancer Therapeutics. <i>ACS Omega</i> , 2022, 7, 25844-25852.	1.6	10
61	Therapeutic targeting of TANK-binding kinase signaling towards anticancer drug development: Challenges and opportunities. <i>International Journal of Biological Macromolecules</i> , 2022, 207, 1022-1037.	3.6	9
62	Cinnamomum zeylanicum Extract and its Bioactive Component Cinnamaldehyde Show Anti-Tumor Effects via Inhibition of Multiple Cellular Pathways. <i>Frontiers in Pharmacology</i> , 0, 13, .	1.6	8
63	Biophysical Insights into Implications of PEG-400 on the Î±-Crystallin Structure: Multispectroscopic and Microscopic Approach. <i>ACS Omega</i> , 2020, 5, 19210-19216.	1.6	7
64	Death-Associated Protein Kinase 3 Inhibitors Identified by Virtual Screening for Drug Discovery in Cancer and Hypertension. <i>OMICS A Journal of Integrative Biology</i> , 2022, 26, 404-413.	1.0	7
65	Structure, function and therapeutic implications of OB-fold proteins: A lesson from past to present. <i>Briefings in Functional Genomics</i> , 2020, 19, 377-389.	1.3	6
66	Targeting metacaspase-3 from <i>Plasmodium falciparum</i> towards antimalarial therapy: A combined approach of <i>in-silico</i> and <i>in-vitro</i> investigation. <i>Journal of Biomolecular Structure and Dynamics</i> , 2021, 39, 421-430.	2.0	6
67	Comparative analysis of web-based programs for single amino acid substitutions in proteins. <i>PLoS ONE</i> , 2022, 17, e0267084.	1.1	6
68	A computational study on active constituents of Habb-ul-aas and Tabasheer as inhibitors of SARS-CoV-2 main protease. <i>Journal of Biomolecular Structure and Dynamics</i> , 2021, , 1-12.	2.0	5
69	Modern Approaches in Synthetic Biology: Genome Editing, Quorum Sensing, and Microbiome Engineering. , 2018, , 189-205.		4
70	Impact of Single Amino Acid Substitutions in Parkinsonism-Associated Deglycase-PARK7 and Their Association with Parkinson's Disease. <i>Journal of Personalized Medicine</i> , 2022, 12, 220.	1.1	4
71	Target-Based Virtual Screening of Natural Compounds Identifies a Potent Antimalarial With Selective Falcipain-2 Inhibitory Activity. <i>Frontiers in Pharmacology</i> , 2022, 13, 850176.	1.6	4
72	Discovering Tuberosin and Villosol as Potent and Selective Inhibitors of AKT1 for Therapeutic Targeting of Oral Squamous Cell Carcinoma. <i>Journal of Personalized Medicine</i> , 2022, 12, 1083.	1.1	3

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73	Investigating the structural features of chromodomain proteins in the human genome and predictive impacts of their mutations in cancers. International Journal of Biological Macromolecules, 2019, 131, 1101-1116.	3.6	2
74	Therapeutic Potential of Polyphenols in Alzheimer's Therapy: Broad-Spectrum and Minimal Side Effects as Key Aspects. , 2021, , 111-133.		2
75	Genome Microbiology for Synthetic Applications. , 2018, , 75-86.		1
76	Implication of Synthetic Biology in Biotherapeutic Engineering. , 2020, , 245-253.		1
77	Role of Genome-Wide Association Studies in Host Genetics: Toward Understanding of Microbiome Association. , 2020, , 37-54.		0
78	ASSESSING THE ASSESSMENT AND EVALUATION PROCESS AT PYP: A CASE STUDY. Indonesian EFL Journal, 2018, 4, 23.	0.3	0
79	Discovery of Novel Drug Targets in Microbial Pathogens Among Hypothetical Proteins: Methods and Significance. , 2019, , 377-391.		0
80	Genome Engineering Tools in Immunotherapy. , 2020, , 73-102.		0
81	Serum and glucocorticoid-regulated kinase 1: A potential target for anticancer therapy. , 2022, , 223-238.		0