

# Rajesh B Patil

## List of Publications by Year in descending order

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32  
papers

588  
citations

840119

11  
h-index

642321

23  
g-index

33  
all docs

33  
docs citations

33  
times ranked

865  
citing authors

#	ARTICLE	IF	CITATIONS
1	Docking Simulations and Primary Assessment of Newly Synthesized Benzene Sulfonamide Pyrazole Oxadiazole Derivatives as Potential Antimicrobial and Antitubercular Agents. Polycyclic Aromatic Compounds, 2023, 43, 1799-1811.	1.4	4
2	Novel $\alpha$ -amylase and $\alpha$ -glucosidase inhibitors from selected Nigerian antidiabetic plants: an <i>in silico</i> approach. Journal of Biomolecular Structure and Dynamics, 2022, 40, 6340-6349.	2.0	11
3	$\alpha$ -Bisabolol Mitigates Colon Inflammation by Stimulating Colon PPAR- $\gamma$ Transcription Factor: In Vivo and In Vitro Study. PPAR Research, 2022, 2022, 1-22.	1.1	7
4	Sars-cov-2 host entry and replication inhibitors from Indian ginseng: an <i>in-silico</i> approach. Journal of Biomolecular Structure and Dynamics, 2021, 39, 4510-4521.	2.0	95
5	Identification of bioactive compounds from <i>Glycyrrhiza glabra</i> as possible inhibitor of SARS-CoV-2 spike glycoprotein and non-structural protein-15: a pharmacoinformatics study. Journal of Biomolecular Structure and Dynamics, 2021, 39, 4686-4700.	2.0	98
6	<i>In-silico</i> investigation of phytochemicals from <i>Asparagus racemosus</i> as plausible antiviral agent in COVID-19. Journal of Biomolecular Structure and Dynamics, 2021, 39, 5033-5047.	2.0	60
7	Computational assessment of saikosaponins as adjuvant treatment for COVID-19: molecular docking, dynamics, and network pharmacology analysis. Molecular Diversity, 2021, 25, 1889-1904.	2.1	25
8	Combination of system biology to probe the anti-viral activity of andrographolide and its derivative against COVID-19. RSC Advances, 2021, 11, 5065-5079.	1.7	28
9	Computational and network pharmacology analysis of bioflavonoids as possible natural antiviral compounds in COVID-19. Informatics in Medicine Unlocked, 2021, 22, 100504.	1.9	36
10	Cocrystals of Apixaban with Improved Solubility and Permeability: Formulation, Physicochemical Characterization, Pharmacokinetic Evaluation, and Computational Studies. Assay and Drug Development Technologies, 2021, 19, 124-138.	0.6	8
11	Computational assessment of select antiviral phytochemicals as potential SARS-Cov-2 main protease inhibitors: molecular dynamics guided ensemble docking and extended molecular dynamics. In Silico Pharmacology, 2021, 9, 44.	1.8	15
12	Computational and network pharmacology studies of Phyllanthus emblica to tackle SARS-CoV-2. Phytomedicine Plus, 2021, 1, 100095.	0.9	12
13	Exploring the therapeutic mechanisms of <i>Cassia glauca</i> in diabetes mellitus through network pharmacology, molecular docking and molecular dynamics. RSC Advances, 2021, 11, 39362-39375.	1.7	16
14	Identification of dual site inhibitors of tankyrase through virtual screening of protein-ligand interaction fingerprint (PLIF)-derived pharmacophore models, molecular dynamics, and ADMET studies. Structural Chemistry, 2020, 31, 769-779.	1.0	1
15	Patent landscape for discovery of promising acyltransferase DGAT and MGAT inhibitors. Expert Opinion on Therapeutic Patents, 2020, 30, 873-896.	2.4	7
16	Pharmacoinformatics approaches to identify potential hits against tetraacyldisaccharide 4 $\epsilon$ -kinase (LpxK) of <i>Pseudomonas aeruginosa</i> . RSC Advances, 2020, 10, 32856-32874.	1.7	3
17	Synthesis and Biological Evaluation of 3,4-Dihydro-2H-benzo[b][1,4]-oxazine-2-carboxylic Acid Derivatives as Antitubercular Agents. Asian Journal of Organic & Medicinal Chemistry, 2020, 5, 138-148.	0.1	0
18	Synthesis and biological activity of structurally diverse phthalazine derivatives: A systematic review. Bioorganic and Medicinal Chemistry, 2019, 27, 3979-3997.	1.4	32

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19	Molecular docking, pharmacophore based virtual screening and molecular dynamics studies towards the identification of potential leads for the management of <i>H. pylori</i> . RSC Advances, 2019, 9, 26176-26208.	1.7	9
20	Comprehensive QSAR studies reveal structural insights into the NR2B subtype selective benzazepine derivatives as N-Methyl-Aspartate receptor antagonists. Journal of Molecular Structure, 2019, 1197, 617-627.	1.8	5
21	Identification of Promising Biofilm Inhibitory and Cytotoxic Quinazolinone Derivatives: Synthesis, Evaluation, Molecular Docking and ADMET Studies. ChemistrySelect, 2019, 4, 3559-3566.	0.7	7
22	Synthesis, biological evaluations and computational studies of N-(3-(2-(7-Chloroquinolin-2-yl)vinyl)) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5 623-630.	1.0	11
23	Preparation, characterization and in vitro evaluation of tablets containing microwave-assisted solid dispersions of apremilast. Polimery W Medycynie, 2019, 48, 17-24.	0.6	10
24	Synthesis, biological evaluation and docking study of some novel isoxazole clubbed 1,3,4-oxadiazoles derivatives. Medicinal Chemistry Research, 2018, 27, 1283-1291.	1.1	19
25	LQTA-R: A new 3D-QSAR methodology applied to a set of DGAT1 inhibitors. Computational Biology and Chemistry, 2018, 74, 123-131.	1.1	11
26	Fungal biofilm inhibition by piperazine-sulphonamide linked Schiff bases: Design, synthesis, and biological evaluation. Archiv Der Pharmazie, 2018, 351, e1700354.	2.1	5
27	Benzene sulfonamide pyrazole thio-oxadiazole hybrid as potential antimicrobial and antitubercular agents. Research on Chemical Intermediates, 2018, 44, 4437-4453.	1.3	9
28	Structural insights of dipeptidyl peptidase-IV inhibitors through molecular dynamics-guided receptor-dependent 4D-QSAR studies. Molecular Diversity, 2018, 22, 575-583.	2.1	5
29	Design, Synthesis and Biological Screening of Novel 1,3,4-Oxadiazoles as Antitubercular Agents. ChemistrySelect, 2018, 3, 13304-13310.	0.7	8
30	Pyrido[1,2-a]pyrimidinones: Ligand-based Design, Synthesis, and Evaluation as an Anti-inflammatory Agent. Journal of Heterocyclic Chemistry, 2017, 54, 3299-3313.	1.4	12
31	4D-QSAR studies of coumarin derivatives as HIV-1 integrase 3'-processing inhibitors. Medicinal Chemistry Research, 2015, 24, 3062-3076.	1.1	8
32	Molecular Dynamics Guided Receptor Independent 4D QSAR Studies of Substituted Coumarins as Anticancer Agents. Current Computer-Aided Drug Design, 2015, 11, 39-50.	0.8	11