

# Tarun Jha

## List of Publications by Year in descending order

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187  
papers

3,849  
citations

147726

31  
h-index

214721

47  
g-index

190  
all docs

190  
docs citations

190  
times ranked

3377  
citing authors

#	ARTICLE	IF	CITATIONS
1	Exploring naphthyl derivatives as SARS-CoV papain-like protease (PLpro) inhibitors and its implications in COVID-19 drug discovery. <i>Molecular Diversity</i> , 2022, 26, 215-228.	2.1	8
2	Synthesis, biological activity, structure activity relationship study and liposomal formulation development of some arylsulfonyl pyroglutamic acid derivatives. <i>Journal of Molecular Structure</i> , 2022, 1248, 131512.	1.8	3
3	How the structural properties of the indole derivatives are important in kinase targeted drug design?: A case study on tyrosine kinase inhibitors. <i>Bioorganic and Medicinal Chemistry</i> , 2022, 53, 116534.	1.4	7
4	Ligand-based quantitative structural assessments of SARS-CoV-2 3CLpro inhibitors: An analysis in light of structure-based multi-molecular modeling evidences. <i>Journal of Molecular Structure</i> , 2022, 1251, 132041.	1.8	12
5	Synthesis, anticancer activity, SAR and binding mode of interaction studies of substituted pentanoic acids: part II. <i>Future Medicinal Chemistry</i> , 2022, 14, 17-34.	1.1	3
6	A critical analysis of urea transporter B inhibitors: molecular fingerprints, pharmacophore features for the development of next-generation diuretics. <i>Molecular Diversity</i> , 2022, 26, 2549-2559.	2.1	1
7	Applying comparative molecular modelling techniques on diverse hydroxamate-based HDAC2 inhibitors: an attempt to identify promising structural features for potent HDAC2 inhibition. <i>SAR and QSAR in Environmental Research</i> , 2022, 33, 1-22.	1.0	8
8	Exploring sodium glucose cotransporter (SGLT2) inhibitors with machine learning approach: A novel hope in anti-diabetes drug discovery. <i>Journal of Molecular Graphics and Modelling</i> , 2022, 111, 108106.	1.3	7
9	Insight into the structural requirements of gelatinases (MMP-2 and MMP-9) inhibitors by multiple validated molecular modelling approaches: Part II. <i>SAR and QSAR in Environmental Research</i> , 2022, 33, 167-192.	1.0	3
10	A fragment-based structural analysis of MMP-2 inhibitors in search of meaningful structural fragments. <i>Computers in Biology and Medicine</i> , 2022, 144, 105360.	3.9	3
11	Binary quantitative activity-activity relationship (QAAR) studies to explore selective HDAC8 inhibitors: In light of mathematical models, DFT-based calculation and molecular dynamic simulation studies. <i>Journal of Molecular Structure</i> , 2022, 1260, 132833.	1.8	4
12	The first report on predictive comparative ligand-based multi-QSAR modeling analysis of 4-pyrimidinone and 2-pyridinone based APJ inhibitors. <i>New Journal of Chemistry</i> , 2022, 46, 11591-11607.	1.4	9
13	Selective inhibition of histone deacetylase 3 by novel hydrazone based small molecules as therapeutic intervention for the treatment of cancer. <i>European Journal of Medicinal Chemistry</i> , 2022, 238, 114470.	2.6	8
14	Chemical-informatics approach to COVID-19 drug discovery: Monte Carlo based QSAR, virtual screening and molecular docking study of some in-house molecules as papain-like protease (PLpro) inhibitors. <i>Journal of Biomolecular Structure and Dynamics</i> , 2021, 39, 4764-4773.	2.0	77
15	Chemical-informatics approach to COVID-19 drug discovery: Exploration of important fragments and data mining based prediction of some hits from natural origins as main protease (Mpro) inhibitors. <i>Journal of Molecular Structure</i> , 2021, 1224, 129026.	1.8	53
16	Structural analysis of arylsulfonamide-based carboxylic acid derivatives: a QSAR study to identify the structural contributors toward their MMP-9 inhibition. <i>Structural Chemistry</i> , 2021, 32, 417-430.	1.0	8
17	Protease targeted COVID-19 drug discovery and its challenges: Insight into viral main protease (Mpro) and papain-like protease (PLpro) inhibitors. <i>Bioorganic and Medicinal Chemistry</i> , 2021, 29, 115860.	1.4	126
18	HDAC6 as privileged target in drug discovery: A perspective. <i>Pharmacological Research</i> , 2021, 163, 105274.	3.1	115

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19	Outline of gelatinase inhibitors as anti-cancer agents: A patent mini-review for 2010-present. <i>European Journal of Medicinal Chemistry</i> , 2021, 213, 113044.	2.6	13
20	Cancer photocytotoxicity and anti-inflammatory response of <i>cis</i> -A <sub>2</sub> B <sub>2</sub> type <i>meso-p</i> -nitrophenyl and <i>p</i> -hydroxyphenyl porphyrin and its zinc(II) complex: a synthetic alternative to the THPP synthon. <i>New Journal of Chemistry</i> , 2021, 45, 2060-2068.	1.4	6
21	Development of decision trees to discriminate HDAC8 inhibitors and non-inhibitors using recursive partitioning. <i>Journal of Biomolecular Structure and Dynamics</i> , 2021, 39, 1-8.	2.0	30
22	Small molecule drug conjugates (SMDCs): an emerging strategy for anticancer drug design and discovery. <i>New Journal of Chemistry</i> , 2021, 45, 5291-5321.	1.4	39
23	First structure-activity relationship analysis of SARS-CoV-2 virus main protease (Mpro) inhibitors: an endeavor on COVID-19 drug discovery. <i>Molecular Diversity</i> , 2021, 25, 1827-1838.	2.1	33
24	Molecular dynamics analysis of phytochemicals from <i>Ageratina adenophora</i> against COVID-19 main protease (Mpro) and human angiotensin-converting enzyme 2 (ACE2). <i>Biocatalysis and Agricultural Biotechnology</i> , 2021, 32, 101924.	1.5	15
25	Protease targeted COVID-19 drug discovery: What we have learned from the past SARS-CoV inhibitors?. <i>European Journal of Medicinal Chemistry</i> , 2021, 215, 113294.	2.6	26
26	Therapies of Hematological Malignancies: An Overview of the Potential Targets and Their Inhibitors. <i>Current Chemical Biology</i> , 2021, 15, 19-49.	0.2	2
27	Robust classification-based molecular modelling of diverse chemical entities as potential SARS-CoV-2 3CL <sup>pro</sup> inhibitors: theoretical justification in light of experimental evidences. <i>SAR and QSAR in Environmental Research</i> , 2021, 32, 473-493.	1.0	9
28	Dissecting Histone Deacetylase 3 in Multiple Disease Conditions: Selective Inhibition as a Promising Therapeutic Strategy. <i>Journal of Medicinal Chemistry</i> , 2021, 64, 8827-8869.	2.9	29
29	Unmasking of crucial structural fragments for coronavirus protease inhibitors and its implications in COVID-19 drug discovery. <i>Journal of Molecular Structure</i> , 2021, 1237, 130366.	1.8	6
30	Insight into the structural requirement of aryl sulphonamide based gelatinases (MMP-2 and MMP-9) inhibitors - Part I: 2D-QSAR, 3D-QSAR topomer CoMFA and Naïve Bayes studies - First report of 3D-QSAR Topomer CoMFA analysis for MMP-9 inhibitors and jointly inhibitors of gelatinases together. <i>SAR and QSAR in Environmental Research</i> , 2021, 32, 655-687.	1.0	4
31	A robust classification-dependent multi-molecular modelling study on some biphenyl sulphonamide based MMP-8 inhibitors. <i>SAR and QSAR in Environmental Research</i> , 2021, 32, 835-861.	1.0	5
32	Synthesis, biological evaluation, and molecular docking analysis of novel linker-less benzamide based potent and selective HDAC3 inhibitors. <i>Bioorganic Chemistry</i> , 2021, 114, 105050.	2.0	14
33	Urea transporter and its specific and nonspecific inhibitors: State of the art and pharmacological perspective. <i>European Journal of Pharmacology</i> , 2021, 911, 174508.	1.7	4
34	Inhibitors of gelatinases (MMP-2 and MMP-9) for the management of hematological malignancies. <i>European Journal of Medicinal Chemistry</i> , 2021, 223, 113623.	2.6	47
35	Quantitative activity-activity relationship (QAAR) driven design to develop hydroxamate derivatives of pentanoic acids as selective HDAC8 inhibitors: synthesis, biological evaluation and binding mode of interaction studies. <i>New Journal of Chemistry</i> , 2021, 45, 17149-17162.	1.4	7
36	Design, synthesis and binding mode of interaction of novel small molecule <i>o</i> -hydroxy benzamides as HDAC3-selective inhibitors with promising antitumor effects in 4T1-Luc breast cancer xenograft model. <i>Bioorganic Chemistry</i> , 2021, 117, 105446.	2.0	13

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37	Ligand-based design of anticancer MMP2 inhibitors: a review. <i>Future Medicinal Chemistry</i> , 2021, 13, 1987-2013.	1.1	9
38	First molecular modelling report on tri-substituted pyrazolines as phosphodiesterase 5 (PDE5) inhibitors through classical and machine learning based multi-QSAR analysis. <i>SAR and QSAR in Environmental Research</i> , 2021, 32, 1-23.	1.0	3
39	Exploration of good and bad structural fingerprints for inhibition of indoleamine-2,3-dioxygenase enzyme in cancer immunotherapy using Monte Carlo optimization and Bayesian classification QSAR modeling. <i>Journal of Biomolecular Structure and Dynamics</i> , 2020, 38, 1-14.	2.0	7
40	Good and bad molecular fingerprints for human rhinovirus 3C protease inhibition: identification, validation, and application in designing of new inhibitors through Monte Carlo-based QSAR study. <i>Journal of Biomolecular Structure and Dynamics</i> , 2020, 38, 66-77.	2.0	34
41	Structural exploration of tetrahydroisoquinoline derivatives as HDAC8 inhibitors through multi-QSAR modeling study. <i>Journal of Biomolecular Structure and Dynamics</i> , 2020, 38, 1551-1564.	2.0	11
42	Essential elements regulating HDAC8 inhibition: a classification based structural analysis and enzyme-inhibitor interaction study of hydroxamate based HDAC8 inhibitors. <i>Journal of Biomolecular Structure and Dynamics</i> , 2020, 38, 5513-5525.	2.0	7
43	Exploration of histone deacetylase 8 inhibitors through classification QSAR study: Part II. <i>Journal of Molecular Structure</i> , 2020, 1204, 127529.	1.8	10
44	Effective anti-aromatase therapy to battle against estrogen-mediated breast cancer: Comparative SAR/QSAR assessment on steroidal aromatase inhibitors. <i>European Journal of Medicinal Chemistry</i> , 2020, 208, 112845.	2.6	9
45	Exploring indole derivatives as myeloid cell leukaemia-1 (Mcl-1) inhibitors with multi-QSAR approach: a novel hope in anti-cancer drug discovery. <i>New Journal of Chemistry</i> , 2020, 44, 17494-17506.	1.4	6
46	Dissecting the Drug Development Strategies Against SARS-CoV-2 Through Diverse Computational Modeling Techniques. <i>Methods in Pharmacology and Toxicology</i> , 2020, , 329-431.	0.1	4
47	<i>meso</i> -Thiophenium Porphyrins and Their Zn(II) Complexes: A New Category of Cationic Photosensitizers. <i>ACS Medicinal Chemistry Letters</i> , 2020, 11, 2041-2047.	1.3	13
48	Photodynamic activity attained through the ruptured $\pi$ -conjugation of pyridyl groups with a porphyrin macrocycle: synthesis and the photophysical and photobiological evaluation of 5-mono-(4-nitrophenyl)-10,15,20-tris-[4-(phenoxy)methyl]pyridine]-porphyrin and its Zn(ii) complex. <i>Photochemical and Photobiological Sciences</i> , 2020, 19, 1776-1789.	1.6	3
49	Folate decorated epigallocatechin-3-gallate (EGCG) loaded PLGA nanoparticles; in-vitro and in-vivo targeting efficacy against MDA-MB-231 tumor xenograft. <i>International Journal of Pharmaceutics</i> , 2020, 585, 119449.	2.6	35
50	Fight against novel coronavirus: A perspective of medicinal chemists. <i>European Journal of Medicinal Chemistry</i> , 2020, 201, 112559.	2.6	58
51	Identification of structural fingerprints for ABCG2 inhibition by using Monte Carlo optimization, Bayesian classification, and structural and physicochemical interpretation (SPCI) analysis. <i>SAR and QSAR in Environmental Research</i> , 2020, 31, 439-455.	1.0	10
52	Matrix metalloproteinase-9 (MMP-9) and its inhibitors in cancer: A minireview. <i>European Journal of Medicinal Chemistry</i> , 2020, 194, 112260.	2.6	270
53	Exploring the structural aspects of ureido-amino acid-based APN inhibitors: a validated comparative multi-QSAR modelling study. <i>SAR and QSAR in Environmental Research</i> , 2020, 31, 325-345.	1.0	5
54	Discriminations of active from inactive HDAC8 inhibitors Part II: Bayesian classification study to find molecular fingerprints. <i>SAR and QSAR in Environmental Research</i> , 2020, 31, 245-260.	1.0	4

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55	Effect of $\hat{\rho}$ coupling on liquid gas phase transition in warm asymmetric nuclear matter. Nuclear Physics A, 2020, 1002, 121974.	0.6	1
56	In silico modelling, identification of crucial molecular fingerprints, and prediction of new possible substrates of human organic cationic transporters 1 and 2. New Journal of Chemistry, 2020, 44, 4129-4143.	1.4	17
57	Histone deacetylase 3 (HDAC3) inhibitors as anticancer agents: A review. European Journal of Medicinal Chemistry, 2020, 192, 112171.	2.6	69
58	Collagenases and gelatinases and their inhibitors as anticancer agents. , 2020, , 265-294.		6
59	Chrysin-loaded PLGA attenuates OVA-induced allergic asthma by modulating TLR/NF- $\hat{\rho}$ B/NLRP3 axis. Nanomedicine: Nanotechnology, Biology, and Medicine, 2020, 30, 102292.	1.7	17
60	An Equation of State for Magnetized Neutron Star Matter and Tidal Deformation in Neutron Star Mergers. Astrophysical Journal, 2020, 900, 49.	1.6	6
61	Designing potential HDAC3 inhibitors to improve memory and learning. Journal of Biomolecular Structure and Dynamics, 2019, 37, 2133-2142.	2.0	21
62	Synthesis, anticancer activity, structure-activity relationship and binding mode of interaction studies of substituted pentanoic acids. Future Medicinal Chemistry, 2019, 11, 1679-1702.	1.1	13
63	Design, synthesis, biological evaluation and molecular docking study of arylcarboxamido piperidine and piperazine-based hydroxamates as potential HDAC8 inhibitors with promising anticancer activity. European Journal of Pharmaceutical Sciences, 2019, 138, 105046.	1.9	32
64	Histone deacetylase 3 inhibitors in learning and memory processes with special emphasis on benzamides. European Journal of Medicinal Chemistry, 2019, 166, 369-380.	2.6	26
65	Intermolecular association of some selected melanin monomers and their optical absorption. Computational and Theoretical Chemistry, 2019, 1151, 43-49.	1.1	6
66	Insight into structural features of phenyltetrazole derivatives as ABCG2 inhibitors for the treatment of multidrug resistance in cancer. SAR and QSAR in Environmental Research, 2019, 30, 457-475.	1.0	19
67	QSAR modelling on a series of arylsulfonamide-based hydroxamates as potent MMP-2 inhibitors. SAR and QSAR in Environmental Research, 2019, 30, 247-263.	1.0	14
68	Dual activity of amphiphilic Zn(II) nitroporphyrin derivatives as HIV-1 entry inhibitors and in cancer photodynamic therapy. European Journal of Medicinal Chemistry, 2019, 174, 66-75.	2.6	29
69	Design of 5-fluorouracil (5-FU) loaded, folate conjugated peptide linked nanoparticles, a potential new drug carrier for selective targeting of tumor cells. MedChemComm, 2019, 10, 559-572.	3.5	19
70	Structural exploration of arylsulfonamide-based ADAM17 inhibitors through validated comparative multi-QSAR modelling studies. Journal of Molecular Structure, 2019, 1185, 128-142.	1.8	9
71	Histone deacetylase 8 (HDAC8) and its inhibitors with selectivity to other isoforms: An overview. European Journal of Medicinal Chemistry, 2019, 164, 214-240.	2.6	72
72	Deconfinement of nonstrange hadronic matter with nucleons and $\hat{\rho}$ baryons to quark matter in neutron stars. International Journal of Modern Physics D, 2019, 28, 1950040.	0.9	9

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73	Reliable structural information for rational design of benzoxazole type potential cholesteryl ester transfer protein (CETP) inhibitors through multiple validated modeling techniques. <i>Journal of Biomolecular Structure and Dynamics</i> , 2019, 37, 4528-4541.	2.0	12
74	Structural refinement and prediction of potential CCR2 antagonists through validated multi-QSAR modeling studies. <i>Journal of Biomolecular Structure and Dynamics</i> , 2019, 37, 75-94.	2.0	11
75	A Review on Camptothecin Analogs with Promising Cytotoxic Profile. <i>Anti-Cancer Agents in Medicinal Chemistry</i> , 2019, 18, 1796-1814.	0.9	20
76	Evaluation of Efficacy of Curcumin along with Lycopene and Piperine in the Management of Oral Submucous Fibrosis. <i>Contemporary Clinical Dentistry</i> , 2019, 10, 531-541.	0.2	10
77	Structure-activity relationships of HDAC8 inhibitors: Non-hydroxamates as anticancer agents. <i>Pharmacological Research</i> , 2018, 131, 128-142.	3.1	32
78	Identification of molecular fingerprints of phenylindole derivatives as cytotoxic agents: a multi-QSAR approach. <i>Structural Chemistry</i> , 2018, 29, 1095-1107.	1.0	9
79	Structural exploration of hydroxyethylamines as HIV-1 protease inhibitors: new features identified. SAR and QSAR in Environmental Research, 2018, 29, 385-408.	1.0	15
80	Design of Aminopeptidase N Inhibitors as Anti-cancer Agents. <i>Journal of Medicinal Chemistry</i> , 2018, 61, 6468-6490.	2.9	83
81	Exploring in house glutamate inhibitors of matrix metalloproteinase-2 through validated robust chemico-biological quantitative approaches. <i>Structural Chemistry</i> , 2018, 29, 285-297.	1.0	16
82	Multiple molecular modelling studies on some derivatives and analogues of glutamic acid as matrix metalloproteinase-2 inhibitors. SAR and QSAR in Environmental Research, 2018, 29, 43-68.	1.0	15
83	Exploring pyrazolo[3,4- <i>d</i> ]pyrimidine phosphodiesterase 1 (PDE1) inhibitors: a predictive approach combining comparative validated multiple molecular modelling techniques. <i>Journal of Biomolecular Structure and Dynamics</i> , 2018, 36, 590-608.	2.0	52
84	Shedding light on designing potential meprin $\beta$ inhibitors through ligand-based robust validated computational approaches: A proposal to chemists!. <i>Journal of Biomolecular Structure and Dynamics</i> , 2018, 36, 3003-3022.	2.0	7
85	Structural exploration for the refinement of anticancer matrix metalloproteinase-2 inhibitor designing approaches through robust validated multi-QSARs. <i>Journal of Molecular Structure</i> , 2018, 1156, 501-515.	1.8	20
86	An integrated QSAR modeling approach to explore the structure–property and selectivity relationships of N-benzoyl-L-biphenylalanines as integrin antagonists. <i>Molecular Diversity</i> , 2018, 22, 129-158.	2.1	10
87	Benzamide porphyrins with directly conjugated and distal pyridyl or pyridinium groups substituted to the porphyrin macrocycles: Study of the photosensitising abilities as inducers of apoptosis in cancer cells under photodynamic conditions. <i>Journal of Photochemistry and Photobiology B: Biology</i> , 2018, 178, 228-236.	1.7	13
88	First Report on the Validated Classification-Based Chemometric Modeling of Human Rhinovirus 3C Protease (HRV 3Cpro) Inhibitors. <i>International Journal of Quantitative Structure-Property Relationships</i> , 2018, 3, 1-20.	1.1	1
89	Selective and nonselective HDAC8 inhibitors: a therapeutic patent review. <i>Pharmaceutical Patent Analyst</i> , 2018, 7, 259-276.	0.4	17
90	Properties of Neutron stars with hyperon cores in parametrized hydrostatic conditions. <i>International Journal of Modern Physics E</i> , 2018, 27, 1850097.	0.4	10

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91	HDAC3 is a potential validated target for cancer: An overview on the benzamide-based selective HDAC3 inhibitors through comparative SAR/QSAR/QAAR approaches. <i>European Journal of Medicinal Chemistry</i> , 2018, 157, 1127-1142.	2.6	48
92	Design, synthesis and biological screening of 2-aminobenzamides as selective HDAC3 inhibitors with promising anticancer effects. <i>European Journal of Pharmaceutical Sciences</i> , 2018, 124, 165-181.	1.9	41
93	Diverse classes of HDAC8 inhibitors: in search of molecular fingerprints that regulate activity. <i>Future Medicinal Chemistry</i> , 2018, 10, 1589-1602.	1.1	30
94	Monte Carlo based modelling approach for designing and predicting cytotoxicity of 2-phenylindole derivatives against breast cancer cell line MCF7. <i>Toxicology in Vitro</i> , 2018, 52, 23-32.	1.1	21
95	Integrin Antagonists: A Special Emphasis on Structural Requirements of N-benzoyl-L-biphenylalanines as $\alpha_5\beta_1$ and $\alpha_4\beta_1$ Antagonists. <i>Current Signal Transduction Therapy</i> , 2018, 13, 105-118.	0.3	1
96	Arylsulfonamides and selectivity of matrix metalloproteinase-2: An overview. <i>European Journal of Medicinal Chemistry</i> , 2017, 129, 72-109.	2.6	55
97	First report on the structural exploration and prediction of new BPTES analogs as glutaminase inhibitors. <i>Journal of Molecular Structure</i> , 2017, 1143, 49-64.	1.8	37
98	Combating breast cancer with non-steroidal aromatase inhibitors (NSAIs): Understanding the chemico-biological interactions through comparative SAR/QSAR study. <i>European Journal of Medicinal Chemistry</i> , 2017, 137, 365-438.	2.6	51
99	Possible anticancer agents: synthesis, pharmacological activity, and molecular modeling studies on some 5-N-Substituted-2-N-(substituted benzenesulphonyl)-L-(+)Glutamines. <i>Medicinal Chemistry Research</i> , 2017, 26, 1437-1458.	1.1	1
100	Insight into the structural requirements of pyrimidine-based phosphodiesterase 10A (PDE10A) inhibitors by multiple validated 3D QSAR approaches. <i>SAR and QSAR in Environmental Research</i> , 2017, 28, 253-273.	1.0	22
101	An integrated ligand-based modelling approach to explore the structure-property relationships of influenza endonuclease inhibitors. <i>Structural Chemistry</i> , 2017, 28, 1663-1678.	1.0	12
102	Hydroxyethylamine derivatives as HIV-1 protease inhibitors: a predictive QSAR modelling study based on Monte Carlo optimization. <i>SAR and QSAR in Environmental Research</i> , 2017, 28, 973-990.	1.0	23
103	A pentanoic acid derivative targeting matrix metalloproteinase-2 (MMP-2) induces apoptosis in a chronic myeloid leukemia cell line. <i>European Journal of Medicinal Chemistry</i> , 2017, 141, 37-50.	2.6	26
104	Homoisoflavonoids as potential antiangiogenic agents for retinal neovascularization. <i>Biomedicine and Pharmacotherapy</i> , 2017, 95, 818-827.	2.5	11
105	Understanding Chemico-Biological Interactions of Glutamate MMP-2 Inhibitors through Rigorous Alignment-Dependent 3D-QSAR Analyses. <i>ChemistrySelect</i> , 2017, 2, 7888-7898.	0.7	12
106	Structure-activity relationships of hydroxamate-based histone deacetylase-8 inhibitors: reality behind anticancer drug discovery. <i>Future Medicinal Chemistry</i> , 2017, 9, 2211-2237.	1.1	31
107	Integrating regression and classification-based QSARs with molecular docking analyses to explore the structure-anti-aromatase activity relationships of letrozole-based analogs. <i>Canadian Journal of Chemistry</i> , 2017, 95, 1285-1295.	0.6	9
108	Structural Insight Into the Viral 3C-Like Protease Inhibitors: Comparative SAR/QSAR Approaches. , 2017, , 317-409.		8

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109	Predictive Quantitative Structure Toxicity Relationship Study on Avian Toxicity of Some Diverse Agrochemical Pesticides by Monte Carlo Method. <i>International Journal of Quantitative Structure-Property Relationships</i> , 2017, 2, 19-34.	1.1	1
110	Is dual inhibition of metalloenzymes HDAC-8 and MMP-2 a potential pharmacological target to combat hematological malignancies?. <i>Pharmacological Research</i> , 2017, 122, 8-19.	3.1	57
111	Designing Potential Antitrypanosomal Thiazol-2-ethylamines through Predictive Regression Based and Classification Based QSAR Analyses. <i>Current Drug Discovery Technologies</i> , 2017, 14, 39-52.	0.6	15
112	Possible Binding Mode Analysis of Pyrazolo-triazole Hybrids as Potential Anticancer Agents through Validated Molecular Docking and 3D-QSAR Modeling Approaches. <i>Letters in Drug Design and Discovery</i> , 2017, 14, 515-527.	0.4	6
113	An Integrated Multi-QSAR Modeling Approach for Designing Knoevenagel- Type Indoles with Enhancing Cytotoxic Profiles. <i>Current Computer-Aided Drug Design</i> , 2017, 13, 336-345.	0.8	6
114	Ligand- and Structure-Based Drug Design of Non-Steroidal Aromatase Inhibitors (NSAIs) in Breast Cancer. , 2017, , 118-186.		0
115	Nitric Oxide Synthase (NOS) Inhibitors in Cancer Angiogenesis. <i>Current Enzyme Inhibition</i> , 2016, 12, 49-66.	0.3	2
116	Robust design of some selective matrix metalloproteinase-2 inhibitors over matrix metalloproteinase-9 through in silico/fragment-based lead identification and de novo lead modification: Syntheses and biological assays. <i>Bioorganic and Medicinal Chemistry</i> , 2016, 24, 4291-4309.	1.4	48
117	First molecular modeling report on novel arylpyrimidine kynurenine monooxygenase inhibitors through multi-QSAR analysis against Huntington's disease: A proposal to chemists!. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2016, 26, 5712-5718.	1.0	41
118	Exploring structural requirements of unconventional Knoevenagel-type indole derivatives as anticancer agents through comparative QSAR modeling approaches. <i>Canadian Journal of Chemistry</i> , 2016, 94, 637-644.	0.6	19
119	A new bisbenzylisoquinoline alkaloid isolated from <i>Thalictrum foliolosum</i> , as a potent inhibitor of DNA topoisomerase IB of <i>Leishmania donovani</i> . <i>FÅ-toterapÅ-Åç</i> , 2016, 109, 25-30.	1.1	30
120	<i>Chenopodium album</i> metabolites act as dual topoisomerase inhibitors and induce apoptosis in the MCF7 cell line. <i>MedChemComm</i> , 2016, 7, 837-844.	3.5	13
121	Insight into the Structural Requirements of Theophylline-Based Aldehyde Dehydrogenase IAL (ALDHIAL) Inhibitors Through Multi-QSAR Modeling and Molecular Docking Approaches. <i>Current Drug Discovery Technologies</i> , 2016, 13, 84-100.	0.6	15
122	Lipid from Infective <i>L. donovani</i> Regulates Acute Myeloid Cell Growth via Mitochondria Dependent MAPK Pathway. <i>PLoS ONE</i> , 2015, 10, e0120509.	1.1	12
123	Anthocephaline, a New Indole Alkaloid and Cadambine, a Potent Inhibitor of DNA Topoisomerase IB of <i>Leishmania donovani</i> (LdTOP1LS), Isolated from <i>Anthocephalus cadamba</i> . <i>Natural Product Communications</i> , 2015, 10, 1934578X1501000.	0.2	23
124	Structural findings of phenylindoles as cytotoxic antimetabolic agents in human breast cancer cell lines through multiple validated QSAR studies. <i>Toxicology in Vitro</i> , 2015, 29, 1392-1404.	1.1	32
125	Leishmanial sphingolipid induces apoptosis in Sarcoma 180 cancer cells through regulation of tumour growth via angiogenic switchover. <i>Tumor Biology</i> , 2015, 36, 3109-3118.	0.8	9
126	Cytotoxic potential of dispirooxindolo/acenaphthoquino andrographolide derivatives against MCF-7 cell line. <i>MedChemComm</i> , 2015, 6, 702-707.	3.5	27



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127	Antineoplastic impact of leishmanial sphingolipid in tumour growth with regulation of angiogenic event and inflammatory response. Apoptosis: an International Journal on Programmed Cell Death, 2015, 20, 869-882.	2.2	7
128	Design of dual MMP-2/HDAC-8 inhibitors by pharmacophore mapping, molecular docking, synthesis and biological activity. RSC Advances, 2015, 5, 72373-72386.	1.7	53
129	Comparative validated molecular modeling of p53-HDM2 inhibitors as antiproliferative agents. European Journal of Medicinal Chemistry, 2015, 90, 860-875.	2.6	24
130	Leishmanial lipid affords protection against oxidative stress induced hepatic injury by regulating inflammatory mediators and confining apoptosis progress. Toxicology Letters, 2015, 232, 499-512.	0.4	15
131	Stepwise development of structure-activity relationship of diverse PARP-1 inhibitors through comparative and validated <i>in silico</i> modeling techniques and molecular dynamics simulation. Journal of Biomolecular Structure and Dynamics, 2015, 33, 1756-1779.	2.0	19
132	Towards the Development of Anticancer Drugs from Andrographolide: Semisynthesis, Bioevaluation, QSAR Analysis and Pharmacokinetic Studies. Current Topics in Medicinal Chemistry, 2015, 15, 1013-1026.	1.0	12
133	Ligand- and Structure-Based Drug Design of Non-Steroidal Aromatase Inhibitors (NSAIs) in Breast Cancer. Advances in Chemical and Materials Engineering Book Series, 2015, , 400-470.	0.2	5
134	Anthocephaline, a new indole alkaloid and cadambine, a potent inhibitor of DNA topoisomerase IB of Leishmania donovani (LdTOP1LS), isolated from Anthocephalus cadamba. Natural Product Communications, 2015, 10, 297-9.	0.2	10
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