

Rajarashi Guha

List of Publications by Year in descending order

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Version: 2024-02-01

58
papers

4,391
citations

186209

28
h-index

161767

54
g-index

62
all docs

62
docs citations

62
times ranked

7497
citing authors

#	ARTICLE	IF	CITATIONS
1	Diversifying cheminformatics. <i>Journal of Cheminformatics</i> , 2022, 14, 25.	2.8	6
2	A High-Throughput Screening Platform Identifies Novel Combination Treatments for Malignant Peripheral Nerve Sheath Tumors. <i>Molecular Cancer Therapeutics</i> , 2022, 21, 1246-1258.	1.9	2
3	What is the role of cheminformatics in a pandemic?. <i>Journal of Cheminformatics</i> , 2021, 13, 16.	2.8	3
4	Reply to "FAIR chemical structure in the <i>Journal of Cheminformatics</i> ". <i>Journal of Cheminformatics</i> , 2021, 13, 49.	2.8	3
5	Brigatinib causes tumor shrinkage in both NF2-deficient meningioma and schwannoma through inhibition of multiple tyrosine kinases but not ALK. <i>PLoS ONE</i> , 2021, 16, e0252048.	1.1	19
6	Learning cheminformatics. <i>Journal of Cheminformatics</i> , 2020, 12, 4.	2.8	7
7	Drugs Targeting Tumor-Initiating Cells Prolong Survival in a Post-Surgery, Post-Chemotherapy Ovarian Cancer Relapse Model. <i>Cancers</i> , 2020, 12, 1645.	1.7	25
8	<i>Journal of Cheminformatics</i> , ORCID, and GitHub. <i>Journal of Cheminformatics</i> , 2019, 11, 44.	2.8	1
9	Scaffold-Based Analytics: Enabling Hit-to-Lead Decisions by Visualizing Chemical Series Linked across Large Datasets. <i>Journal of Chemical Information and Modeling</i> , 2019, 59, 4880-4892.	2.5	8
10	High-Throughput Screening for Drug Combinations. <i>Methods in Molecular Biology</i> , 2019, 1939, 11-35.	0.4	10
11	Implementing cheminformatics. <i>Journal of Cheminformatics</i> , 2019, 11, 12.	2.8	0
12	Therapeutic strategies for diffuse midline glioma from high-throughput combination drug screening. <i>Science Translational Medicine</i> , 2019, 11, .	5.8	129
13	The Rise and Fall of a Scaffold: A Trend Analysis of Scaffolds in the Medicinal Chemistry Literature. <i>Journal of Medicinal Chemistry</i> , 2018, 61, 4688-4703.	2.9	41
14	Canvass: A Crowd-Sourced, Natural-Product Screening Library for Exploring Biological Space. <i>ACS Central Science</i> , 2018, 4, 1727-1741.	5.3	32
15	Using Machine Learning to Predict Synergistic Antimalarial Compound Combinations With Novel Structures. <i>Frontiers in Pharmacology</i> , 2018, 9, 1096.	1.6	27
16	MEK inhibition induces MYOG and remodels super-enhancers in RAS-driven rhabdomyosarcoma. <i>Science Translational Medicine</i> , 2018, 10, .	5.8	104
17	Identification of Combinations of Approved Drugs With Synergistic Activity Against Ebola Virus in Cell Cultures. <i>Journal of Infectious Diseases</i> , 2018, 218, S672-S678.	1.9	49
18	A systematic and prospectively validated approach for identifying synergistic drug combinations against malaria. <i>Malaria Journal</i> , 2018, 17, 160.	0.8	19

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19	Pharmacological and genomic profiling of neurofibromatosis type 1 plexiform neurofibroma-derived schwann cells. <i>Scientific Data</i> , 2018, 5, 180106.	2.4	20
20	Pharos: Collating protein information to shed light on the druggable genome. <i>Nucleic Acids Research</i> , 2017, 45, D995-D1002.	6.5	271
21	PAX3 ^{hi} FOXO1 Establishes Myogenic Super Enhancers and Confers BET Bromodomain Vulnerability. <i>Cancer Discovery</i> , 2017, 7, 884-899.	7.7	221
22	Helping to improve the practice of cheminformatics. <i>Journal of Cheminformatics</i> , 2017, 9, 40.	2.8	3
23	The Chemistry Development Kit (CDK) v2.0: atom typing, depiction, molecular formulas, and substructure searching. <i>Journal of Cheminformatics</i> , 2017, 9, 33.	2.8	275
24	Modelling of compound combination effects and applications to efficacy and toxicity: state-of-the-art, challenges and perspectives. <i>Drug Discovery Today</i> , 2016, 21, 225-238.	3.2	162
25	Augmented efficacy of brentuximab vedotin combined with ruxolitinib and/or Navitoclax in a murine model of human Hodgkin ⁺ lymphoma. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 1624-1629.	3.3	38
26	High-throughput matrix screening identifies synergistic and antagonistic antimalarial drug combinations. <i>Scientific Reports</i> , 2015, 5, 13891.	1.6	92
27	BioAssay Research Database (BARD): chemical biology and probe-development enabled by structured metadata and result types. <i>Nucleic Acids Research</i> , 2015, 43, D1163-D1170.	6.5	20
28	Synergy Maps: exploring compound combinations using network-based visualization. <i>Journal of Cheminformatics</i> , 2015, 7, 36.	2.8	32
29	High-throughput combinatorial screening identifies drugs that cooperate with ibrutinib to kill activated B-cell ⁺ like diffuse large B-cell lymphoma cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 2349-2354.	3.3	355
30	Blockade of oncogenic I ^h B kinase activity in diffuse large B-cell lymphoma by bromodomain and extraterminal domain protein inhibitors. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 11365-11370.	3.3	166
31	On the validity versus utility of activity landscapes: are all activity cliffs statistically significant?. <i>Journal of Cheminformatics</i> , 2014, 6, 11.	2.8	9
32	What are we ⁺ tweeting ⁺ about obesity? Mapping tweets with topic modeling and Geographic Information System. <i>Cartography and Geographic Information Science</i> , 2013, 40, 90-102.	1.4	171
33	A Survey of Quantitative Descriptions of Molecular Structure. <i>Current Topics in Medicinal Chemistry</i> , 2012, 12, 1946-1956.	1.0	48
34	Cheminformatics. <i>Communications of the ACM</i> , 2012, 55, 65-75.	3.3	21
35	Exploring Uncharted Territories: Predicting Activity Cliffs in Structure ⁺ Activity Landscapes. <i>Journal of Chemical Information and Modeling</i> , 2012, 52, 2181-2191.	2.5	33
36	Exploring structure ⁺ activity data using the landscape paradigm. <i>Wiley Interdisciplinary Reviews: Computational Molecular Science</i> , 2012, 2, 829-841.	6.2	34

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37	Chemical Genomic Profiling for Antimalarial Therapies, Response Signatures, and Molecular Targets. <i>Science</i> , 2011, 333, 724-729.	6.0	130
38	Using a neural network for mining interpretable relationships of West Nile risk factors. <i>Social Science and Medicine</i> , 2011, 72, 418-429.	1.8	13
39	Exploratory analysis of kinetic solubility measurements of a small molecule library. <i>Bioorganic and Medicinal Chemistry</i> , 2011, 19, 4127-4134.	1.4	27
40	Open Data, Open Source and Open Standards in chemistry: The Blue Obelisk five years on. <i>Journal of Cheminformatics</i> , 2011, 3, 37.	2.8	63
41	The Ups and Downs of Structure-Activity Landscapes. <i>Methods in Molecular Biology</i> , 2010, 672, 101-117.	0.4	28
42	Use of genetic algorithm and neural network approaches for risk factor selection: A case study of West Nile virus dynamics in an urban environment. <i>Computers, Environment and Urban Systems</i> , 2010, 34, 189-203.	3.3	14
43	Advances in Cheminformatics Methodologies and Infrastructure to Support the Data Mining of Large, Heterogeneous Chemical Datasets. <i>Current Computer-Aided Drug Design</i> , 2010, 6, 50-67.	0.8	28
44	GOAssay: from Gene Ontology to Assays Identifiers - Towards Automatic Functional Annotation of PubChem BioAssays. <i>Nature Precedings</i> , 2009, , .	0.1	1
45	PubChem as a Source of Polypharmacology. <i>Journal of Chemical Information and Modeling</i> , 2009, 49, 2044-2055.	2.5	130
46	Utilizing high throughput screening data for predictive toxicology models: protocols and application to MLSCN assays. <i>Journal of Computer-Aided Molecular Design</i> , 2008, 22, 367-384.	1.3	43
47	On the interpretation and interpretability of quantitative structure-activity relationship models. <i>Journal of Computer-Aided Molecular Design</i> , 2008, 22, 857-871.	1.3	70
48	Structure-Activity Landscape Index: Identifying and Quantifying Activity Cliffs. <i>Journal of Chemical Information and Modeling</i> , 2008, 48, 646-658.	2.5	281
49	Assessing How Well a Modeling Protocol Captures a Structure-Activity Landscape. <i>Journal of Chemical Information and Modeling</i> , 2008, 48, 1716-1728.	2.5	84
50	SQMD: Architecture for Scalable, Distributed Database System Built on Virtual Private Servers. , 2008, , .		2
51	Open Notebook Science - Falcipain-2 Preliminary Results. <i>Nature Precedings</i> , 2008, , .	0.1	1
52	The CombiUgi Project and Closing the Open Science Loop. <i>Nature Precedings</i> , 2007, , .	0.1	0
53	Chemical Informatics Functionality in R. <i>Journal of Statistical Software</i> , 2007, 18, .	1.8	189
54	The Blue Obelisk-Interoperability in Chemical Informatics. <i>Journal of Chemical Information and Modeling</i> , 2006, 46, 991-998.	2.5	366

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55	Scalable Partitioning and Exploration of Chemical Spaces Using Geometric Hashing. Journal of Chemical Information and Modeling, 2006, 46, 321-333.	2.5	22
56	Recent Developments of the Chemistry Development Kit (CDK) - An Open-Source Java Library for Chemo- and Bioinformatics. Current Pharmaceutical Design, 2006, 12, 2111-2120.	0.9	418
57	Determining the Validity of a QSAR Model ? A Classification Approach.. ChemInform, 2005, 36, no.	0.1	0
58	Development of QSAR Models to Predict and Interpret the Biological Activity of Artemisinin Analogues.. ChemInform, 2004, 35, no.	0.1	0