

Florian Auer

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4943109/publications.pdf>

Version: 2024-02-01

10
papers

899
citations

1684188

5
h-index

1720034

7
g-index

15
all docs

15
docs citations

15
times ranked

1510
citing authors

#	ARTICLE	IF	CITATIONS
1	RCXâ€™ an R package adapting the Cytoscape Exchange format for biological networks. <i>Bioinformatics Advances</i> , 2022, 2, .	2.4	1
2	Implementation of Gene Expression Profiles in the HL7 FHIR Standard. <i>Studies in Health Technology and Informatics</i> , 2022, , .	0.3	0
3	MISeval: A Metric Library for Medical Image Segmentation Evaluation. <i>Studies in Health Technology and Informatics</i> , 2022, , .	0.3	6
4	Perspective on Code Submission and Automated Evaluation Platforms for University Teaching. <i>Studies in Health Technology and Informatics</i> , 2022, , .	0.3	0
5	Data-dependent visualization of biological networks in the web-browser with NDExEdit. <i>PLoS Computational Biology</i> , 2022, 18, e1010205.	3.2	1
6	Adaptation of HL7 FHIR for the Exchange of Patientsâ€™™ Gene Expression Profiles. <i>Studies in Health Technology and Informatics</i> , 2022, , .	0.3	0
7	Explaining decisions of graph convolutional neural networks: patient-specific molecular subnetworks responsible for metastasis prediction in breast cancer. <i>Genome Medicine</i> , 2021, 13, 42.	8.2	48
8	ndexrâ€™ an R package to interface with the network data exchange. <i>Bioinformatics</i> , 2018, 34, 716-717.	4.1	11
9	Homology-based inference sets the bar high for protein function prediction. <i>BMC Bioinformatics</i> , 2013, 14, S7.	2.6	38
10	A large-scale evaluation of computational protein function prediction. <i>Nature Methods</i> , 2013, 10, 221-227.	19.0	789