

Ritambhara Singh

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5995524/publications.pdf>

Version: 2024-02-01

13
papers

514
citations

1478458

6
h-index

1372553

10
g-index

19
all docs

19
docs citations

19
times ranked

612
citing authors

#	ARTICLE	IF	CITATIONS
1	An automated COVID-19 triage pipeline using artificial intelligence based on chest radiographs and clinical data. <i>Npj Digital Medicine</i> , 2022, 5, 5.	10.9	22
2	SCOT: Single-Cell Multi-Omics Alignment with Optimal Transport. <i>Journal of Computational Biology</i> , 2022, 29, 3-18.	1.6	40
3	Single-Cell Multiomics Integration by SCOT. <i>Journal of Computational Biology</i> , 2022, 29, 19-22.	1.6	3
4	Integrating Long-Range Regulatory Interactions to Predict Gene Expression Using Graph Convolutional Networks. <i>Journal of Computational Biology</i> , 2022, 29, 409-424.	1.6	9
5	A pan-tissue DNA-methylation epigenetic clock based on deep learning. , 2022, 8, .		27
6	Unsupervised Integration of Single-Cell Multi-omics Datasets with Disproportionate Cell-Type Representation. <i>Lecture Notes in Computer Science</i> , 2022, , 3-19.	1.3	4
7	TIMEOR: a web-based tool to uncover temporal regulatory mechanisms from multi-omics data. <i>Nucleic Acids Research</i> , 2021, 49, W641-W653.	14.5	9
8	Single-cell landscape of nuclear configuration and gene expression during stem cell differentiation and X inactivation. <i>Genome Biology</i> , 2021, 22, 279.	8.8	11
9	Machine learning for profile prediction in genomics. <i>Current Opinion in Chemical Biology</i> , 2021, 65, 35-41.	6.1	11
10	Unsupervised manifold alignment for single-cell multi-omics data. , 2020, 2020, 1-10.		33
11	FastSK: fast sequence analysis with gapped string kernels. <i>Bioinformatics</i> , 2020, 36, i857-i865.	4.1	5
12	Jointly Embedding Multiple Single-Cell Omics Measurements. , 2019, 143, .		25
13	DeepChrome: deep-learning for predicting gene expression from histone modifications. <i>Bioinformatics</i> , 2016, 32, i639-i648.	4.1	224