

Jason Liu

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1051840/publications.pdf>

Version: 2024-02-01

13
papers

215
citations

1478505

6
h-index

1125743

13
g-index

13
all docs

13
docs citations

13
times ranked

441
citing authors

#	ARTICLE	IF	CITATIONS
1	An integrative ENCODE resource for cancer genomics. <i>Nature Communications</i> , 2020, 11, 3696.	12.8	95
2	STARRPeaker: uniform processing and accurate identification of STARR-seq active regions. <i>Genome Biology</i> , 2020, 21, 298.	8.8	36
3	Establishing a Global Standard for Wearable Devices in Sport and Fitness: Perspectives from the New England Chapter of the American College of Sports Medicine Members. <i>Current Sports Medicine Reports</i> , 2020, 19, 45-49.	1.2	18
4	Epigenome-based splicing prediction using a recurrent neural network. <i>PLoS Computational Biology</i> , 2020, 16, e1008006.	3.2	16
5	Establishing a Global Standard for Wearable Devices in Sport and Exercise Medicine: Perspectives from Academic and Industry Stakeholders. <i>Sports Medicine</i> , 2021, 51, 2237-2250.	6.5	12
6	RADAR: annotation and prioritization of variants in the post-transcriptional regulome of RNA-binding proteins. <i>Genome Biology</i> , 2020, 21, 151.	8.8	9
7	DECODE: a deep-learning framework for decoding enhancers and refining boundaries with large-scale functional assays. <i>Bioinformatics</i> , 2021, 37, i280-i288.	4.1	8
8	Bayesian structural time series for biomedical sensor data: A flexible modeling framework for evaluating interventions. <i>PLoS Computational Biology</i> , 2021, 17, e1009303.	3.2	8
9	DiNeR: a Differential graphical model for analysis of co-regulation Network Rewiring. <i>BMC Bioinformatics</i> , 2020, 21, 281.	2.6	5
10	Cross-platform transcriptomic profiling of the response to recombinant human erythropoietin. <i>Scientific Reports</i> , 2021, 11, 21705.	3.3	5
11	GRAM: A GeneRALized Model to predict the molecular effect of a non-coding variant in a cell-type specific manner. <i>PLoS Genetics</i> , 2019, 15, e1007860.	3.5	1
12	NIMBus: a negative binomial regression based Integrative Method for mutation Burden Analysis. <i>BMC Bioinformatics</i> , 2020, 21, 474.	2.6	1
13	Gene Tracer: a smart, interactive, voice-controlled Alexa skill For gene information retrieval and browsing, mutation annotation and network visualization. <i>Bioinformatics</i> , 2021, 37, 2998-3000.	4.1	1