

Sai Zhang

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/12147406/publications.pdf>

Version: 2024-02-01

21
papers

811
citations

759233

12
h-index

794594

19
g-index

27
all docs

27
docs citations

27
times ranked

1225
citing authors

#	ARTICLE	IF	CITATIONS
1	A deep learning framework for modeling structural features of RNA-binding protein targets. <i>Nucleic Acids Research</i> , 2016, 44, e32-e32.	14.5	213
2	TITER: predicting translation initiation sites by deep learning. <i>Bioinformatics</i> , 2017, 33, i234-i242.	4.1	83
3	Gene-Environment Interaction in the Era of Precision Medicine. <i>Cell</i> , 2019, 177, 38-44.	28.9	73
4	Decoding the Genomics of Abdominal Aortic Aneurysm. <i>Cell</i> , 2018, 174, 1361-1372.e10.	28.9	68
5	Physical exercise is a risk factor for amyotrophic lateral sclerosis: Convergent evidence from Mendelian randomisation, transcriptomics and risk genotypes. <i>EBioMedicine</i> , 2021, 68, 103397.	6.1	65
6	Analysis of Ribosome Stalling and Translation Elongation Dynamics by Deep Learning. <i>Cell Systems</i> , 2017, 5, 212-220.e6.	6.2	58
7	Genome-wide identification of the genetic basis of amyotrophic lateral sclerosis. <i>Neuron</i> , 2022, 110, 992-1008.e11.	8.1	51
8	DeepHINT: understanding HIV-1 integration via deep learning with attention. <i>Bioinformatics</i> , 2019, 35, 1660-1667.	4.1	41
9	A review of Mendelian randomization in amyotrophic lateral sclerosis. <i>Brain</i> , 2022, 145, 832-842.	7.6	29
10	Rare Variant Burden Analysis within Enhancers Identifies CAV1 as an ALS Risk Gene. <i>Cell Reports</i> , 2020, 33, 108456.	6.4	24
11	A deep boosting based approach for capturing the sequence binding preferences of RNA-binding proteins from high-throughput CLIP-seq data. <i>Nucleic Acids Research</i> , 2017, 45, e129-e129.	14.5	19
12	Membrane lipid raft homeostasis is directly linked to neurodegeneration. <i>Essays in Biochemistry</i> , 2021, 65, 999-1011.	4.7	15
13	Precision medicine in women with epilepsy: The challenge, systematic review, and future direction. <i>Epilepsy and Behavior</i> , 2021, 118, 107928.	1.7	13
14	Advances in the genetic classification of amyotrophic lateral sclerosis. <i>Current Opinion in Neurology</i> , 2021, 34, 756-764.	3.6	12
15	Unbiased metabolome screen leads to personalized medicine strategy for amyotrophic lateral sclerosis. <i>Brain Communications</i> , 2022, 4, fcac069.	3.3	10
16	Multiomic analysis reveals cell-type-specific molecular determinants of COVID-19 severity. <i>Cell Systems</i> , 2022, 13, 598-614.e6.	6.2	10
17	Characterizing RNA Pseudouridylation by Convolutional Neural Networks. <i>Genomics, Proteomics and Bioinformatics</i> , 2021, 19, 815-833.	6.9	5
18	Elastic restricted Boltzmann machines for cancer data analysis. <i>Quantitative Biology</i> , 2017, 5, 159-172.	0.5	4

#	ARTICLE	IF	CITATIONS
19	ROSE: A Deep Learning Based Framework for Predicting Ribosome Stalling. SSRN Electronic Journal, 0, , .	0.4	2
20	DeepRibSt: a multi-feature convolutional neural network for predicting ribosome stalling. Multimedia Tools and Applications, 2021, 80, 17239-17255.	3.9	1
21	Genome-Wide Identification of the Genetic Basis of Amyotrophic Lateral Sclerosis. SSRN Electronic Journal, 0, , .	0.4	1