

Seonwoo Min

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3965135/publications.pdf>

Version: 2024-02-01

14
papers

1,905
citations

759233

12
h-index

1058476

14
g-index

14
all docs

14
docs citations

14
times ranked

2768
citing authors

#	ARTICLE	IF	CITATIONS
1	Deep learning in bioinformatics. <i>Briefings in Bioinformatics</i> , 2017, 18, bbw068.	6.5	865
2	Deep learning improves prediction of CRISPRâ€“Cpf1 guide RNA activity. <i>Nature Biotechnology</i> , 2018, 36, 239-241.	17.5	252
3	Predicting the efficiency of prime editing guide RNAs in human cells. <i>Nature Biotechnology</i> , 2021, 39, 198-206.	17.5	160
4	Prediction of the sequence-specific cleavage activity of Cas9 variants. <i>Nature Biotechnology</i> , 2020, 38, 1328-1336.	17.5	133
5	SpCas9 activity prediction by DeepSpCas9, a deep learningâ€“based model with high generalization performance. <i>Science Advances</i> , 2019, 5, eaax9249.	10.3	130
6	High-throughput analysis of the activities of xCas9, SpCas9-NG and SpCas9 at matched and mismatched target sequences in human cells. <i>Nature Biomedical Engineering</i> , 2020, 4, 111-124.	22.5	98
7	Sequence-specific prediction of the efficiencies of adenine and cytosine base editors. <i>Nature Biotechnology</i> , 2020, 38, 1037-1043.	17.5	73
8	Learned Embeddings from Deep Learning to Visualize and Predict Protein Sets. <i>Current Protocols</i> , 2021, 1, e113.	2.9	61
9	Generation of a more efficient prime editor 2 by addition of the Rad51 DNA-binding domain. <i>Nature Communications</i> , 2021, 12, 5617.	12.8	47
10	Recording of elapsed time and temporal information about biological events using Cas9. <i>Cell</i> , 2021, 184, 1047-1063.e23.	28.9	29
11	Pre-Training of Deep Bidirectional Protein Sequence Representations With Structural Information. <i>IEEE Access</i> , 2021, 9, 123912-123926.	4.2	20
12	Learning-Based Instantaneous Drowsiness Detection Using Wired and Wireless Electroencephalography. <i>IEEE Access</i> , 2019, 7, 146390-146402.	4.2	14
13	TargetNet: functional microRNA target prediction with deep neural networks. <i>Bioinformatics</i> , 2022, 38, 671-677.	4.1	12
14	Protein transfer learning improves identification of heat shock protein families. <i>PLoS ONE</i> , 2021, 16, e0251865.	2.5	11