

Kemal Eren

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11507508/publications.pdf>

Version: 2024-02-01

11
papers

1,418
citations

1163117

8
h-index

1281871

11
g-index

13
all docs

13
docs citations

13
times ranked

2701
citing authors

#	ARTICLE	IF	CITATIONS
1	Gene-Wide Identification of Episodic Selection. <i>Molecular Biology and Evolution</i> , 2015, 32, 1365-1371.	8.9	493
2	Antibody 10-1074 suppresses viremia in HIV-1-infected individuals. <i>Nature Medicine</i> , 2017, 23, 185-191.	30.7	399
3	A comparative analysis of biclustering algorithms for gene expression data. <i>Briefings in Bioinformatics</i> , 2013, 14, 279-292.	6.5	196
4	Early Antibody Lineage Diversification and Independent Limb Maturation Lead to Broad HIV-1 Neutralization Targeting the Env High-Mannose Patch. <i>Immunity</i> , 2016, 44, 1215-1226.	14.3	138
5	HIV Envelope Glycoform Heterogeneity and Localized Diversity Govern the Initiation and Maturation of a V2 Apex Broadly Neutralizing Antibody Lineage. <i>Immunity</i> , 2017, 47, 990-1003.e9.	14.3	90
6	Long-read amplicon denoising. <i>Nucleic Acids Research</i> , 2019, 47, e104-e104.	14.5	31
7	Rapid Sequencing of Complete <i>env</i> Genes from Primary HIV-1 Samples. <i>Virus Evolution</i> , 2016, 2, vew018.	4.9	30
8	HIV Superinfection Drives De Novo Antibody Responses and Not Neutralization Breadth. <i>Cell Host and Microbe</i> , 2018, 24, 593-599.e3.	11.0	24
9	Full-Length Envelope Analyzer (FLEA): A tool for longitudinal analysis of viral amplicons. <i>PLoS Computational Biology</i> , 2018, 14, e1006498.	3.2	5
10	RIFRAF: a frame-resolving consensus algorithm. <i>Bioinformatics</i> , 2018, 34, 3817-3824.	4.1	4
11	Querying Co-regulated Genes on Diverse Gene Expression Datasets Via Biclustering. <i>Methods in Molecular Biology</i> , 2015, 1375, 55-74.	0.9	1