

Ferhat Alkan

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/818135/publications.pdf>

Version: 2024-02-01

13
papers

481
citations

1163117

8
h-index

1125743

13
g-index

15
all docs

15
docs citations

15
times ranked

681
citing authors

#	ARTICLE	IF	CITATIONS
1	CRISPR-Cas9 off-targeting assessment with nucleic acid duplex energy parameters. <i>Genome Biology</i> , 2018, 19, 177.	8.8	105
2	Rlsearch2: suffix array-based large-scale prediction of RNA-RNA interactions and siRNA off-targets. <i>Nucleic Acids Research</i> , 2017, 45, gkw1325.	14.5	75
3	BEAMS: backbone extraction and merge strategy for the global many-to-many alignment of multiple PPI networks. <i>Bioinformatics</i> , 2014, 30, 531-539.	4.1	63
4	Good guide, bad guide: spacer sequence-dependent cleavage efficiency of Cas12a. <i>Nucleic Acids Research</i> , 2020, 48, 3228-3243.	14.5	62
5	RAIN: RNA-protein Association and Interaction Networks. <i>Database: the Journal of Biological Databases and Curation</i> , 2017, 2017, baw167.	3.0	53
6	Glucocorticoid receptor triggers a reversible drug-tolerant dormancy state with acquired therapeutic vulnerabilities in lung cancer. <i>Nature Communications</i> , 2021, 12, 4360.	12.8	35
7	CRISPR/Cas9 gRNA activity depends on free energy changes and on the target PAM context. <i>Nature Communications</i> , 2022, 13, .	12.8	31
8	Joint Profiling of miRNAs and mRNAs Reveals miRNA Mediated Gene Regulation in the Göttingen Minipig Obesity Model. <i>PLoS ONE</i> , 2016, 11, e0167285.	2.5	14
9	BSGatlas: a unified <i>Bacillus subtilis</i> genome and transcriptome annotation atlas with enhanced information access. <i>Microbial Genomics</i> , 2021, 7, .	2.0	12
10	SiPAN: simultaneous prediction and alignment of protein-protein interaction networks. <i>Bioinformatics</i> , 2015, 31, 2356-2363.	4.1	9
11	Identifying ribosome heterogeneity using ribosome profiling. <i>Nucleic Acids Research</i> , 2022, 50, e95-e95.	14.5	7
12	RedNemo: topology-based PPI network reconstruction via repeated diffusion with neighborhood modifications. <i>Bioinformatics</i> , 2017, 33, 537-544.	4.1	6
13	<i>Ribo-ODDR</i> : oligo design pipeline for experiment-specific rRNA depletion in Ribo-seq. <i>Bioinformatics</i> , 2021, 37, 2659-2667.	4.1	5