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List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1019427/publications.pdf>

Version: 2024-02-01

9
papers

341
citations

1478505

6
h-index

1474206

9
g-index

12
all docs

12
docs citations

12
times ranked

461
citing authors

#	ARTICLE	IF	CITATIONS
1	snoDB: an interactive database of human snoRNA sequences, abundance and interactions. <i>Nucleic Acids Research</i> , 2020, 48, D220-D225.	14.5	83
2	Simultaneous sequencing of coding and noncoding RNA reveals a human transcriptome dominated by a small number of highly expressed noncoding genes. <i>Rna</i> , 2018, 24, 950-965.	3.5	61
3	Protein coding genes as hosts for noncoding RNA expression. <i>Seminars in Cell and Developmental Biology</i> , 2018, 75, 3-12.	5.0	52
4	Identification of discrete classes of small nucleolar RNA featuring different ends and RNA binding protein dependency. <i>Nucleic Acids Research</i> , 2014, 42, 10073-10085.	14.5	46
5	Handling multi-mapped reads in RNA-seq. <i>Computational and Structural Biotechnology Journal</i> , 2020, 18, 1569-1576.	4.1	44
6	CoCo: RNA-seq read assignment correction for nested genes and multimapped reads. <i>Bioinformatics</i> , 2019, 35, 5039-5047.	4.1	28
7	SnoRNA copy regulation affects family size, genomic location and family abundance levels. <i>BMC Genomics</i> , 2021, 22, 414.	2.8	12
8	High-grade ovarian cancer associated H/ACA snoRNAs promote cancer cell proliferation and survival. <i>NAR Cancer</i> , 2022, 4, zcab050.	3.1	10
9	The snoGloBe interaction predictor reveals a broad spectrum of C/D snoRNA RNA targets. <i>Nucleic Acids Research</i> , 2022, 50, 6067-6083.	14.5	4