Guanqun Zheng

List of Publications by Year in Descending Order

Source: https://exaly.com/author-pdf/12012351/guanqun-zheng-publications-by-year.pdf

Version: 2024-04-09

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

6,811 21 20 20 h-index g-index citations papers 8,701 21 22.1 5.37 L-index ext. citations avg, IF ext. papers

#	Paper	IF	Citations
20	Determination of tRNA aminoacylation levels by high-throughput sequencing. <i>Nucleic Acids Research</i> , 2017 , 45, e133	20.1	33
19	Selective Enzymatic Demethylation of N2,N2-Dimethylguanosine in RNA and Its Application in High-Throughput tRNA Sequencing. <i>Angewandte Chemie</i> , 2017 , 129, 5099-5102	3.6	3
18	Selective Enzymatic Demethylation of N ,N -Dimethylguanosine in RNA and Its Application in High-Throughput tRNA Sequencing. <i>Angewandte Chemie - International Edition</i> , 2017 , 56, 5017-5020	16.4	30
17	ALKBH1-Mediated tRNA Demethylation Regulates Translation. Cell, 2016, 167, 816-828.e16	56.2	197
16	The dynamic N(1)-methyladenosine methylome in eukaryotic messenger RNA. <i>Nature</i> , 2016 , 530, 441-6	50.4	523
15	tRNA base methylation identification and quantification via high-throughput sequencing. <i>Rna</i> , 2016 , 22, 1771-1784	5.8	100
14	N(6)-methyladenosine-dependent RNA structural switches regulate RNA-protein interactions. <i>Nature</i> , 2015 , 518, 560-4	50.4	988
13	Kinetic gating mechanism of DNA damage recognition by Rad4/XPC. <i>Nature Communications</i> , 2015 , 6, 5849	17.4	59
12	Efficient and quantitative high-throughput tRNA sequencing. <i>Nature Methods</i> , 2015 , 12, 835-837	21.6	291
11	Synthesis of a FTO inhibitor with anticonvulsant activity. ACS Chemical Neuroscience, 2014, 5, 658-65	5.7	59
10	Dynamics of spontaneous flipping of a mismatched base in DNA duplex. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 8043-8	11.5	64
9	Nucleic acid oxidation in DNA damage repair and epigenetics. Chemical Reviews, 2014, 114, 4602-20	68.1	63
8	Unique features of the m6A methylome in Arabidopsis thaliana. <i>Nature Communications</i> , 2014 , 5, 5630	17.4	239
7	Crystal structure of the RNA demethylase ALKBH5 from zebrafish. FEBS Letters, 2014, 588, 892-8	3.8	41
6	Probing N6-methyladenosine RNA modification status at single nucleotide resolution in mRNA and long noncoding RNA. <i>Rna</i> , 2013 , 19, 1848-56	5.8	320
5	ALKBH5 is a mammalian RNA demethylase that impacts RNA metabolism and mouse fertility. <i>Molecular Cell</i> , 2013 , 49, 18-29	17.6	1627
4	Sprouts of RNA epigenetics: the discovery of mammalian RNA demethylases. RNA Biology, 2013, 10, 915	5 zβ 8	69

LIST OF PUBLICATIONS

Chemical Biology, 2011, 7, 885-7

3

2	Iron-catalysed oxidation intermediates captured in a DNA repair dioxygenase. <i>Nature</i> , 2010 , 468, 330-3	50.4	109
	Structure determination of DNA methylation lesions N1-meA and N3-meC in duplex DNA using a	• • •	26

 ${\sf N6-methylade} no sine in nuclear \ {\sf RNA} is a major substrate of the obesity-associated \ {\sf FTO}. \ {\it Nature}$

cross-linked protein-DNA system. Nucleic Acids Research, 2010, 38, 4415-25

11.7 1937

20.1 36