Guan-Zheng Luo

List of Publications by Year in Descending Order

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

53	4,657	29	54
papers	citations	h-index	g-index
54	6,355 ext. citations	15.4	5.37
ext. papers		avg, IF	L-index

#	Paper	IF	Citations
53	The RNA mA reader YTHDC1 silences retrotransposons and guards ES cell identity. <i>Nature</i> , 2021 , 591, 322-326	50.4	45
52	Targeted RNA N -Methyladenosine Demethylation Controls Cell Fate Transition in Human Pluripotent Stem Cells. <i>Advanced Science</i> , 2021 , 8, e2003902	13.6	7
51	The Impact of Microbiome and Microbiota-Derived Sodium Butyrate on Transcriptome and Metabolome Revealed by Multi-Omics Analysis. <i>Metabolites</i> , 2021 , 11,	5.6	3
50	Mapping single-nucleotide mA by mA-REF-seq. <i>Methods</i> , 2021 ,	4.6	1
49	Long noncoding RNA sponges mmu-miR-139-5p to modulate functions in mouse ESCs and embryos. <i>RNA Biology</i> , 2021 , 18, 875-887	4.8	4
48	RNA mA Modification Functions in Larval Development and Caste Differentiation in Honeybee (Apis mellifera). <i>Cell Reports</i> , 2021 , 34, 108580	10.6	5
47	Targeted genetic screening in bacteria with a Cas12k-guided transposase. <i>Cell Reports</i> , 2021 , 36, 10963	5 10.6	1
46	Establishment of transposase-assisted low-input mA sequencing technique. <i>Journal of Genetics and Genomics</i> , 2021 , 48, 1036-1039	4	1
45	Systematic calibration of epitranscriptomic maps using a synthetic modification-free RNA library. <i>Nature Methods</i> , 2021 , 18, 1213-1222	21.6	4
44	Mapping and editing of nucleic acid modifications. <i>Computational and Structural Biotechnology Journal</i> , 2020 , 18, 661-667	6.8	7
43	Peroxisome Elevation Induces Stem Cell Differentiation and Intestinal Epithelial Repair. <i>Developmental Cell</i> , 2020 , 53, 169-184.e11	10.2	11
42	Keth-seq for transcriptome-wide RNA structure mapping. <i>Nature Chemical Biology</i> , 2020 , 16, 489-492	11.7	31
41	Crystal structure of the yeast heterodimeric ADAT2/3 deaminase. <i>BMC Biology</i> , 2020 , 18, 189	7.3	4
40	Acute Deletion of METTL14 in ECells of Adult Mice Results in Glucose Intolerance. <i>Endocrinology</i> , 2019 , 160, 2388-2394	4.8	10
39	RNA mA methylation regulates the epithelial mesenchymal transition of cancer cells and translation of Snail. <i>Nature Communications</i> , 2019 , 10, 2065	17.4	234
38	METTL14 is essential for Etell survival and insulin secretion. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2019 , 1865, 2138-2148	6.9	19
37	Transcriptome-wide Mapping of Internal N-Methylguanosine Methylome in Mammalian mRNA. <i>Molecular Cell</i> , 2019 , 74, 1304-1316.e8	17.6	133

36	Single-base mapping of mA by an antibody-independent method. <i>Science Advances</i> , 2019 , 5, eaax0250	14.3	128
35	Transcriptome-wide reprogramming of N-methyladenosine modification by the mouse microbiome. <i>Cell Research</i> , 2019 , 29, 167-170	24.7	19
34	Transfer RNA demethylase ALKBH3 promotes cancer progression via induction of tRNA-derived small RNAs. <i>Nucleic Acids Research</i> , 2019 , 47, 2533-2545	20.1	108
33	VIRMA mediates preferential mA mRNA methylation in 3WTR and near stop codon and associates with alternative polyadenylation. <i>Cell Discovery</i> , 2018 , 4, 10	22.3	332
32	N-methyldeoxyadenosine directs nucleosome positioning in Tetrahymena DNA. <i>Genome Biology</i> , 2018 , 19, 200	18.3	26
31	Mapping and characterizing N6-methyladenine in eukaryotic genomes using single-molecule real-time sequencing. <i>Genome Research</i> , 2018 , 28, 1067-1078	9.7	48
30	Ythdc2 is an N-methyladenosine binding protein that regulates mammalian spermatogenesis. <i>Cell Research</i> , 2017 , 27, 1115-1127	24.7	404
29	YTHDC1 mediates nuclear export of N-methyladenosine methylated mRNAs. <i>ELife</i> , 2017 , 6,	8.9	452
28	Author response: YTHDC1 mediates nuclear export of N6-methyladenosine methylated mRNAs 2017 ,		6
27	DNA N-methyladenine in metazoans: functional epigenetic mark or bystander?. <i>Nature Structural and Molecular Biology</i> , 2017 , 24, 503-506	17.6	54
26	ALKBH1-Mediated tRNA Demethylation Regulates Translation. Cell, 2016, 167, 816-828.e16	56.2	197
25	Characterization of eukaryotic DNA N(6)-methyladenine by a highly sensitive restriction enzyme-assisted sequencing. <i>Nature Communications</i> , 2016 , 7, 11301	17.4	62
24	Abundant DNA 6mA methylation during early embryogenesis of zebrafish and pig. <i>Nature Communications</i> , 2016 , 7, 13052	17.4	141
23	Structure and mechanism of the essential two-component signal-transduction system WalKR in Staphylococcus aureus. <i>Nature Communications</i> , 2016 , 7, 11000	17.4	21
22	Ubiquitously expressed genes participate in cell-specific functions via alternative promoter usage. <i>EMBO Reports</i> , 2016 , 17, 1304-13	6.5	14
21	High-Resolution Mapping of NEMethyladenosine in Transcriptome and Genome Using a Photo-Crosslinking-Assisted Strategy. <i>Methods in Enzymology</i> , 2015 , 560, 161-85	1.7	16
20	Widespread occurrence of N6-methyladenosine in bacterial mRNA. <i>Nucleic Acids Research</i> , 2015 , 43, 65	57:67	117
19	Durable pluripotency and haploidy in epiblast stem cells derived from haploid embryonic stem cells in vitro. <i>Journal of Molecular Cell Biology</i> , 2015 , 7, 326-37	6.3	16

N6-methyldeoxyadenosine marks active transcription start sites in Chlamydomonas. Cell, 2015, 161, 879-202 316 18 DNA N(6)-methyladenine: a new epigenetic mark in eukaryotes?. Nature Reviews Molecular Cell 48.7 17 157 Biology, 2015, 16, 705-10 High-Resolution N6-Methyladenosine (m6A) Map Using Photo-Crosslinking-Assisted m6A 16 3.6 26 Sequencing. *Angewandte Chemie*, **2015**, 127, 1607-1610 High-resolution N(6) -methyladenosine (m(6) A) map using photo-crosslinking-assisted m(6) A 16.4 15 249 sequencing. Angewandte Chemie - International Edition, 2015, 54, 1587-90 Genetic modification and screening in rat using haploid embryonic stem cells. Cell Stem Cell, 2014, 18 14 71 14.404-14 Unique features of the m6A methylome in Arabidopsis thaliana. Nature Communications, 2014, 5, 5630 17.4 13 239 MicroRNA-323-3p regulates the activity of polycomb repressive complex 2 (PRC2) via targeting the mRNA of embryonic ectoderm development (Eed) gene in mouse embryonic stem cells. Journal of 6 12 5.4 Biological Chemistry, 2013, 288, 23659-65 Parthenogenetic haploid embryonic stem cells produce fertile mice. Cell Research, 2013, 23, 1330-3 11 28 24.7 In vivo suppression of microRNA-24 prevents the transition toward decompensated hypertrophy in 10 15.7 71 aortic-constricted mice. Circulation Research, 2013, 112, 601-5 miR-9 and miR-140-5p target FoxP2 and are regulated as a function of the social context of singing 6.6 41 9 behavior in zebra finches. Journal of Neuroscience, 2013, 33, 16510-21 8 Three-dimensional culture may promote cell reprogramming. Organogenesis, 2013, 9, 118-20 1.7 7 Identification of a small molecule 1,4-bis-[4-(3-phenoxy-propoxy)-but-2-ynyl]-piperazine as a novel 8 inhibitor of the transcription factor p53. Acta Pharmacologica Sinica, 2013, 34, 805-10 6 Mir-24 regulates junctophilin-2 expression in cardiomyocytes. Circulation Research, 2012, 111, 837-41 15.7 74 Androgenetic haploid embryonic stem cells produce live transgenic mice. Nature, 2012, 490, 407-11 129 Dynamics of brassinosteroid response modulated by negative regulator LIC in rice. PLoS Genetics, 6 95 2012, 8, e1002686 Upregulation of a disintegrin and metalloproteinase with thrombospondin motifs-7 by miR-29 repression mediates vascular smooth muscle calcification. Arteriosclerosis, Thrombosis, and Vascular 96 9.4 Biology, 2012, 32, 2580-8 Activation of the imprinted Dlk1-Dio3 region correlates with pluripotency levels of mouse stem 218 5.4 cells. Journal of Biological Chemistry, 2010, 285, 19483-90 BC10, a DUF266-containing and Golgi-located type II membrane protein, is required for cell-wall 88 6.9 biosynthesis in rice (Oryza sativa L.). Plant Journal, 2009, 57, 446-62