

MODOMICS: a database of RNA modification pathways.

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Challenges with Simulating Modified RNA: Insights into Role and Reciprocity of Experimental and Computational Approaches. <i>Genes</i> , 2022, 13, 540.	2.4	4
3	The 2022 <i>Nucleic Acids Research</i> database issue and the online molecular biology database collection. <i>Nucleic Acids Research</i> , 2022, 50, D1-D10.	14.5	50
4	Decoding pseudouridine: an emerging target for therapeutic development. <i>Trends in Pharmacological Sciences</i> , 2022, 43, 522-535.	8.7	32
5	m5CRegpred: Epitranscriptome Target Prediction of 5-Methylcytosine (m5C) Regulators Based on Sequencing Features. <i>Genes</i> , 2022, 13, 677.	2.4	9
7	The plant epitranscriptome: revisiting pseudouridine and 2-O-methyl RNA modifications. <i>Plant Biotechnology Journal</i> , 2022, 20, 1241-1256.	8.3	10
9	Developmental mRNA m5C landscape and regulatory innovations of massive m5C modification of maternal mRNAs in animals. <i>Nature Communications</i> , 2022, 13, 2484.	12.8	24
10	The tRNA regulome in neurodevelopmental and neuropsychiatric disease. <i>Molecular Psychiatry</i> , 2022, 27, 3204-3213.	7.9	9
11	Research Progress of RNA Methylation Modification in Colorectal Cancer. <i>Frontiers in Pharmacology</i> , 2022, 13, .	3.5	3
12	Mass spectrometry profiling analysis enables the identification of new modifications in ribosomal RNA. <i>Chinese Chemical Letters</i> , 2023, 34, 107531.	9.0	12
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14	m1A RNA Modification in Gene Expression Regulation. <i>Genes</i> , 2022, 13, 910.	2.4	28
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19	Rapid Determination of RNA Modifications in Consensus Motifs by Nuclease Protection with Ion-Tagged Oligonucleotide Probes and Matrix-Assisted Laser Desorption Ionization Mass Spectrometry. <i>Genes</i> , 2022, 13, 1008.	2.4	1
20	On demand CRISPR-mediated RNA N6-methyladenosine editing. <i>Genes and Diseases</i> , 2022, 9, 1389-1390.	3.4	3
21	Research Progress of RNA m5C Methylation Modification and Its Detection Technology in Tumors. <i>Advances in Clinical Medicine</i> , 2022, 12, 5407-5414.	0.0	0
22	Construction of a Novel Prognostic Model in Lung Adenocarcinoma Based on 7-Methylguanosine-Related Gene Signatures. <i>Frontiers in Oncology</i> , 0, 12, .	2.8	5

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25	Analysis of queuosine and 2-thio tRNA modifications by high throughput sequencing. <i>Nucleic Acids Research</i> , 2022, 50, e99-e99.	14.5	8
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