

Todd M Lowe

List of Publications by Year in descending order

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Version: 2024-02-01

36
papers

23,158
citations

201674

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37
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docs citations

45
times ranked

23997
citing authors

#	ARTICLE	IF	CITATIONS
1	Eukaryotic tRNA sequences present conserved and amino acid-specific structural signatures. <i>Nucleic Acids Research</i> , 2022, 50, 4100-4112.	14.5	6
2	Distinct Stressâ€Dependent Signatures of Cellular and Extracellular tRNAâ€Derived Small RNAs. <i>Advanced Science</i> , 2022, 9, e2200829.	11.2	19
3	R2DT is a framework for predicting and visualising RNA secondary structure using templates. <i>Nature Communications</i> , 2021, 12, 3494.	12.8	58
4	tRNAscan-SE 2.0: improved detection and functional classification of transfer RNA genes. <i>Nucleic Acids Research</i> , 2021, 49, 9077-9096.	14.5	569
5	Predicting transfer RNA gene activity from sequence and genome context. <i>Genome Research</i> , 2020, 30, 85-94.	5.5	22
6	A guide to naming human nonâ€coding RNA genes. <i>EMBO Journal</i> , 2020, 39, e103777.	7.8	77
7	Distinct Modified Nucleosides in tRNA ^{Trp} from the Hyperthermophilic Archaeon <i>Thermococcus kodakarensis</i> and Requirement of tRNA m ² G10/m ² G10 Methyltransferase (Archaeal Trm11) for Survival at High Temperatures. <i>Journal of Bacteriology</i> , 2019, 201,...	2.2	15
8	Matching tRNA modifications in humans to their known and predicted enzymes. <i>Nucleic Acids Research</i> , 2019, 47, 2143-2159.	14.5	116
9	tRNAviz: explore and visualize tRNA sequence features. <i>Nucleic Acids Research</i> , 2019, 47, W542-W547.	14.5	17
10	tRNAscan-SE: Searching for tRNA Genes in Genomic Sequences. <i>Methods in Molecular Biology</i> , 2019, 1962, 1-14.	0.9	1,023
11	RNAcentral: a hub of information for non-coding RNA sequences. <i>Nucleic Acids Research</i> , 2019, 47, D221-D229.	14.5	153
12	Stress response of a marine ammonia-oxidizing archaeon informs physiological status of environmental populations. <i>ISME Journal</i> , 2018, 12, 508-519.	9.8	82
13	Methylation guide RNA evolution in archaea: structure, function and genomic organization of 110 C/D box sRNA families across six <i>Pyrobaculum</i> species. <i>Nucleic Acids Research</i> , 2018, 46, 5678-5691.	14.5	7
14	Transfer RNA genes experience exceptionally elevated mutation rates. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 8996-9001.	7.1	40
15	High-Throughput Small RNA Sequencing Enhanced by AlkB-Facilitated RNA de-Methylation (ARM-Seq). <i>Methods in Molecular Biology</i> , 2017, 1562, 231-243.	0.9	17
16	RNAcentral: a comprehensive database of non-coding RNA sequences. <i>Nucleic Acids Research</i> , 2017, 45, D128-D134.	14.5	174
17	tRNAscan-SE On-line: integrating search and context for analysis of transfer RNA genes. <i>Nucleic Acids Research</i> , 2016, 44, W54-W57.	14.5	2,219
18	Small RNA Modifications: Integral to Function and Disease. <i>Trends in Molecular Medicine</i> , 2016, 22, 1025-1034.	6.7	90

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19	GtRNAdb 2.0: an expanded database of transfer RNA genes identified in complete and draft genomes. <i>Nucleic Acids Research</i> , 2016, 44, D184-D189.	14.5	776
20	ARM-seq: AlkB-facilitated RNA methylation sequencing reveals a complex landscape of modified tRNA fragments. <i>Nature Methods</i> , 2015, 12, 879-884.	19.0	350
21	RNAcentral: an international database of ncRNA sequences. <i>Nucleic Acids Research</i> , 2015, 43, D123-D129.	14.5	103
22	C/D box sRNA-guided 2â€²-O-methylation patterns of archaeal rRNA molecules. <i>BMC Genomics</i> , 2015, 16, 632.	2.8	35
23	Draft De Novo Transcriptome of the Rat Kangaroo Potorous tridactylus as a Tool for Cell Biology. <i>PLoS ONE</i> , 2015, 10, e0134738.	2.5	18
24	IscR Is Essential for Yersinia pseudotuberculosis Type III Secretion and Virulence. <i>PLoS Pathogens</i> , 2014, 10, e1004194.	4.7	53
25	Reclassification of <i>Thermoproteus neutrophilus</i> Stetter and Zillig 1989 as <i>Pyrobaculum neutrophilum</i> comb. nov. based on phylogenetic analysis. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 751-754.	1.7	21
26	Small nucleolar RNAs and RNA-guided post-transcriptional modification. <i>Essays in Biochemistry</i> , 2013, 54, 53-77.	4.7	69
27	Complete genome sequence of <i>Pyrobaculum oguniense</i> . <i>Standards in Genomic Sciences</i> , 2012, 6, 336-345.	1.5	10
28	Diversity of Antisense and Other Non-Coding RNAs in Archaea Revealed by Comparative Small RNA Sequencing in Four <i>Pyrobaculum</i> Species. <i>Frontiers in Microbiology</i> , 2012, 3, 231.	3.5	46
29	The UCSC Archaeal Genome Browser: 2012 update. <i>Nucleic Acids Research</i> , 2012, 40, D646-D652.	14.5	89
30	Discovery of permuted and recently split transfer RNAs in Archaea. <i>Genome Biology</i> , 2011, 12, R38.	8.8	58
31	GtRNAdb: a database of transfer RNA genes detected in genomic sequence. <i>Nucleic Acids Research</i> , 2009, 37, D93-D97.	14.5	782
32	The tRNAscan-SE, snoscan and snoGPS web servers for the detection of tRNAs and snoRNAs. <i>Nucleic Acids Research</i> , 2005, 33, W686-W689.	14.5	2,090
33	A guided tour: small RNA function in Archaea. <i>Molecular Microbiology</i> , 2001, 40, 509-519.	2.5	128
34	Homologs of Small Nucleolar RNAs in Archaea. <i>Science</i> , 2000, 288, 517-522.	12.6	324
35	tRNAscan-SE: A Program for Improved Detection of Transfer RNA Genes in Genomic Sequence. <i>Nucleic Acids Research</i> , 1997, 25, 955-964.	14.5	9,417
36	tRNAscan-SE: A Program for Improved Detection of Transfer RNA Genes in Genomic Sequence. <i>Nucleic Acids Research</i> , 1997, 25, 0955-964.	14.5	3,970