

Mario MÃ¶rl

List of Publications by Year in descending order

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Version: 2024-02-01

65
papers

3,119
citations

218677

26
h-index

168389

53
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72
all docs

72
docs citations

72
times ranked

3441
citing authors

#	ARTICLE	IF	CITATIONS
1	Synthetic Riboswitches for the Analysis of tRNA Processing by eukaryotic RNase P Enzymes. <i>Rna</i> , 2022, , rna.078814.121.	3.5	3
2	RNA Design Principles for Riboswitches that Regulate RNase P-Mediated tRNA Processing. <i>Methods in Molecular Biology</i> , 2022, , 179-202.	0.9	1
3	CCA-Addition Gone Wild: Unusual Occurrence and Phylogeny of Four Different tRNA Nucleotidyltransferases in <i>Acanthamoeba castellanii</i> . <i>Molecular Biology and Evolution</i> , 2021, 38, 1006-1017.	8.9	0
4	Beyond Plug and Pray: Context Sensitivity and <i>in silico</i> Design of Artificial Neomycin Riboswitches. <i>RNA Biology</i> , 2021, 18, 457-467.	3.1	6
5	Ligand-dependent tRNA processing by a rationally designed RNase P riboswitch. <i>Nucleic Acids Research</i> , 2021, 49, 1784-1800.	14.5	4
6	Crystallization and Structural Determination of an Enzyme:Substrate Complex by Serial Crystallography in a Versatile Microfluidic Chip. <i>Journal of Visualized Experiments</i> , 2021, , .	0.3	0
7	Changes of the tRNA Modification Pattern during the Development of <i>Dictyostelium discoideum</i> . <i>Non-coding RNA</i> , 2021, 7, 32.	2.6	1
8	CCA-addition in the cold: Structural characterization of the psychrophilic CCA-adding enzyme from the permafrost bacterium <i>Planococcus halocryophilus</i> . <i>Computational and Structural Biotechnology Journal</i> , 2021, 19, 5845-5855.	4.1	2
9	LOTTE-seq (Long hairpin oligonucleotide based tRNA high-throughput sequencing): specific selection of tRNAs with 3â€™-CCA end for high-throughput sequencing. <i>RNA Biology</i> , 2020, 17, 23-32.	3.1	22
10	Unusual Occurrence of Two Bona-Fide CCA-Adding Enzymes in <i>Dictyostelium discoideum</i> . <i>International Journal of Molecular Sciences</i> , 2020, 21, 5210.	4.1	4
11	Adaptation of the <i>Romanomermis culicivorax</i> CCA-Adding Enzyme to Miniaturized Armless tRNA Substrates. <i>International Journal of Molecular Sciences</i> , 2020, 21, 9047.	4.1	6
12	Divergent Evolution of Eukaryotic CC- and A-Adding Enzymes. <i>International Journal of Molecular Sciences</i> , 2020, 21, 462.	4.1	5
13	Monitoring the Production of High Diffraction-Quality Crystals of Two Enzymes in Real Time Using In Situ Dynamic Light Scattering. <i>Crystals</i> , 2020, 10, 65.	2.2	3
14	Evolving methods for rational de novo design of functional RNA molecules. <i>Methods</i> , 2019, 161, 54-63.	3.8	6
15	Dual expression of CCA-adding enzyme and RNase T in <i>Escherichia coli</i> generates a distinct cca growth phenotype with diverse applications. <i>Nucleic Acids Research</i> , 2019, 47, 3631-3639.	14.5	7
16	A Temporal Order in 5â€™- and 3â€™- Processing of Eukaryotic tRNAHis. <i>International Journal of Molecular Sciences</i> , 2019, 20, 1384.	4.1	3
17	Post-Transcriptional Regulation of tRNA Pools To Govern the Central Dogma: A Perspective. <i>Biochemistry</i> , 2019, 58, 299-304.	2.5	1
18	A simple and versatile microfluidic device for efficient biomacromolecule crystallization and structural analysis by serial crystallography. <i>IUCr</i> , 2019, 6, 454-464.	2.2	23

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19	Designer tRNAs for efficient incorporation of non-canonical amino acids by the pyrrolysine system in mammalian cells. <i>Nucleic Acids Research</i> , 2018, 46, 1-10.	14.5	170
20	Accurate mapping of tRNA reads. <i>Bioinformatics</i> , 2018, 34, 1116-1124.	4.1	33
21	A tRNA's fate is decided at its 3' end: Collaborative actions of CCA-adding enzyme and RNases involved in tRNA processing and degradation. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2018, 1861, 433-441.	1.9	41
22	Cold adaptation of tRNA nucleotidyltransferases: A tradeoff in activity, stability and fidelity. <i>RNA Biology</i> , 2018, 15, 144-155.	3.1	24
23	Examining tRNA 3' ends in <i>Escherichia coli</i> : teamwork between CCA-adding enzyme, RNase T, and RNase R. <i>Rna</i> , 2018, 24, 361-370.	3.5	20
24	Combining crystallography methods to produce diffraction-quality crystals of a psychrophilic tRNA-maturation enzyme. <i>Acta Crystallographica Section F, Structural Biology Communications</i> , 2018, 74, 747-753.	0.8	8
25	Small but large enough: structural properties of armless mitochondrial tRNAs from the nematode <i>Romanomermis culicivorax</i> . <i>Nucleic Acids Research</i> , 2018, 46, 9170-9180.	14.5	35
26	Applicability of a computational design approach for synthetic riboswitches. <i>Nucleic Acids Research</i> , 2017, 45, gkw1267.	14.5	52
27	Synthetic Riboswitches: From Plug and Pray toward Plug and Play. <i>Biochemistry</i> , 2017, 56, 1181-1198.	2.5	82
28	tRNA Modifications: Impact on Structure and Thermal Adaptation. <i>Biomolecules</i> , 2017, 7, 35.	4.0	241
29	Design of Artificial Riboswitches as Biosensors. <i>Sensors</i> , 2017, 17, 1990.	3.8	50
30	Genotyping bacterial and fungal pathogens using sequence variation in the gene for the CCA-adding enzyme. <i>BMC Microbiology</i> , 2016, 16, 47.	3.3	5
31	The CCA-adding enzyme: A central scrutinizer in tRNA quality control. <i>BioEssays</i> , 2015, 37, 975-982.	2.5	35
32	The identity of the discriminator base has an impact on CCA addition. <i>Nucleic Acids Research</i> , 2015, 43, 5617-5629.	14.5	22
33	The ancestor of modern Holozoa acquired the CCA-adding enzyme from Alphaproteobacteria by horizontal gene transfer. <i>Nucleic Acids Research</i> , 2015, 43, 6739-6746.	14.5	14
34	Design of Transcription Regulating Riboswitches. <i>Methods in Enzymology</i> , 2015, 550, 1-22.	1.0	8
35	Design criteria for synthetic riboswitches acting on transcription. <i>RNA Biology</i> , 2015, 12, 221-231.	3.1	41
36	Domain movements during CCA-addition: A new function for motif C in the catalytic core of the human tRNA nucleotidyltransferases. <i>RNA Biology</i> , 2015, 12, 435-446.	3.1	14

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37	From End to End: tRNA Editing at 5'- and 3'-Terminal Positions. <i>International Journal of Molecular Sciences</i> , 2014, 15, 23975-23998.	4.1	27
38	Biological evidence for the world's smallest tRNAs. <i>Biochimie</i> , 2014, 100, 151-158.	2.6	57
39	Reversible and Rapid Transfer-RNA Deactivation as a Mechanism of Translational Repression in Stress. <i>PLoS Genetics</i> , 2013, 9, e1003767.	3.5	94
40	De novo design of a synthetic riboswitch that regulates transcription termination. <i>Nucleic Acids Research</i> , 2013, 41, 2541-2551.	14.5	163
41	The TRAMP Complex Shows tRNA Editing Activity in <i>S. cerevisiae</i> . <i>Molecular Biology and Evolution</i> , 2012, 29, 1451-1459.	8.9	6
42	A new mitochondrial point mutation in the transfer RNALys gene associated with progressive external ophthalmoplegia with impaired respiratory regulation. <i>Journal of the Neurological Sciences</i> , 2012, 316, 108-111.	0.6	9
43	An inhibitory C-terminal region dictates the specificity of A-adding enzymes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 21040-21045.	7.1	20
44	tRNA nucleotidyltransferases: ancient catalysts with an unusual mechanism of polymerization. <i>Cellular and Molecular Life Sciences</i> , 2010, 67, 1447-1463.	5.4	62
45	tRNA nucleotidyltransferases: Highly unusual RNA polymerases with vital functions. <i>FEBS Letters</i> , 2010, 584, 297-302.	2.8	26
46	Unusual evolution of a catalytic core element in CCA-adding enzymes. <i>Nucleic Acids Research</i> , 2010, 38, 4436-4447.	14.5	19
47	tRNAdb 2009: compilation of tRNA sequences and tRNA genes. <i>Nucleic Acids Research</i> , 2009, 37, D159-D162.	14.5	751
48	A comparative analysis of CCA-adding enzymes from human and <i>E. coli</i> : Differences in CCA addition and tRNA 3'-end repair. <i>Biochimie</i> , 2008, 90, 762-772.	2.6	42
49	Evolution of tRNA nucleotidyltransferases: A small deletion generated CC-adding enzymes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 7953-7958.	7.1	42
50	A comparative analysis of two conserved motifs in bacterial poly(A) polymerase and CCA-adding enzyme. <i>Nucleic Acids Research</i> , 2008, 36, 5212-5220.	14.5	25
51	A Splice Variant of the Human CCA-adding Enzyme with Modified Activity. <i>Journal of Molecular Biology</i> , 2007, 366, 1258-1265.	4.2	13
52	Hfq stimulates the activity of the CCA-adding enzyme. <i>BMC Molecular Biology</i> , 2007, 8, 92.	3.0	22
53	Is yeast on its way to evolving tRNA editing?. <i>EMBO Reports</i> , 2005, 6, 367-372.	4.5	10
54	Exchange of Regions between Bacterial Poly(A) Polymerase and the CCA-Adding Enzyme Generates Altered Specificities. <i>Molecular Cell</i> , 2004, 15, 389-398.	9.7	46

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55	A Pathogenesis-associated Mutation in Human Mitochondrial tRNA ^{Leu} (UUR) Leads to Reduced 3'-End Processing and CCA Addition. <i>Journal of Molecular Biology</i> , 2004, 337, 535-544.	4.2	60
56	Mitochondrial tRNA editing. <i>Topics in Current Genetics</i> , 2004, , 81-96.	0.7	1
57	Crystal Structure of the Human CCA-adding Enzyme: Insights into Template-independent Polymerization. <i>Journal of Molecular Biology</i> , 2003, 328, 985-994.	4.2	71
58	A universal method to produce in vitro transcripts with homogeneous 3' ends. <i>Nucleic Acids Research</i> , 2002, 30, 56e-56.	14.5	140
59	The final cut. <i>EMBO Reports</i> , 2001, 2, 17-20.	4.5	97
60	This Is the End: Processing, Editing and Repair at the tRNA 3-Terminus. <i>Biological Chemistry</i> , 2001, 382, 1147-56.	2.5	73
61	Evidence for Import of a Lysyl-tRNA into Marsupial Mitochondria. <i>Molecular Biology of the Cell</i> , 2001, 12, 2688-2698.	2.1	82
62	Repair of tRNAs in metazoan mitochondria. <i>Nucleic Acids Research</i> , 2000, 28, 2043-2048.	14.5	34
63	Processing and Editing of Overlapping tRNAs in Human Mitochondria. <i>Journal of Biological Chemistry</i> , 1998, 273, 31977-31984.	3.4	46
64	C to U editing and modifications during the maturation of the mitochondrial tRNA ^{ASP} in marsupials. <i>Nucleic Acids Research</i> , 1995, 23, 3380-3384.	14.5	68
65	Production of RNAs with Homogeneous 5' and 3' Ends. , 0, , 22-35.		13