

# Richard J Maraia

## List of Publications by Year in descending order

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96  
papers

5,056  
citations

61977

43  
h-index

102480

66  
g-index

104  
all docs

104  
docs citations

104  
times ranked

6367  
citing authors

#	ARTICLE	IF	CITATIONS
1	A Role for TFIIIC Transcription Factor Complex in Genome Organization. <i>Cell</i> , 2006, 125, 859-872.	28.9	275
2	A La-Related Protein Modulates 7SK snRNP Integrity to Suppress P-TEFb-Dependent Transcriptional Elongation and Tumorigenesis. <i>Molecular Cell</i> , 2008, 29, 588-599.	9.7	223
3	BCR/ABL activates mdm2 mRNA translation via the La antigen. <i>Cancer Cell</i> , 2003, 3, 145-160.	16.8	199
4	Transcriptional regulation and transpositional selection of active SINE sequences. <i>Current Opinion in Genetics and Development</i> , 1992, 2, 874-882.	3.3	163
5	Alu transcripts: cytoplasmic localisation and regulation by DNA methylation. <i>Nucleic Acids Research</i> , 1994, 22, 1087-1095.	14.5	152
6	Conserved and divergent features of the structure and function of La and La-related proteins (LARPs). <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2010, 1799, 365-378.	1.9	128
7	Defective i6A37 Modification of Mitochondrial and Cytosolic tRNAs Results from Pathogenic Mutations in TRIT1 and Its Substrate tRNA. <i>PLoS Genetics</i> , 2014, 10, e1004424.	3.5	112
8	Transcription termination factor La is also an initiation factor for RNA polymerase III.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1996, 93, 3383-3387.	7.1	111
9	Phosphorylation of the Human La Antigen on Serine 366 Can Regulate Recycling of RNA Polymerase III Transcription Complexes. <i>Cell</i> , 1997, 88, 707-715.	28.9	111
10	5â€™ Processing of tRNA Precursors Can Be Modulated by the Human La Antigen Phosphoprotein. <i>Molecular and Cellular Biology</i> , 1998, 18, 3201-3211.	2.3	110
11	Recognition of Nascent RNA by the Human La Antigen: Conserved and Divergent Features of Structure and Function. <i>Molecular and Cellular Biology</i> , 2001, 21, 367-379.	2.3	109
12	Differential Phosphorylation and Subcellular Localization of La RNPs Associated with Precursor tRNAs and Translation-Related mRNAs. <i>Molecular Cell</i> , 2003, 12, 1301-1307.	9.7	109
13	3â€™ processing of eukaryotic precursor tRNAs. <i>Wiley Interdisciplinary Reviews RNA</i> , 2011, 2, 362-375.	6.4	107
14	The <sc>La</sc> and related <sc>RNA</sc>-binding proteins (LARPs): structures, functions, and evolving perspectives. <i>Wiley Interdisciplinary Reviews RNA</i> , 2017, 8, e1430.	6.4	99
15	Transcription termination by the eukaryotic RNA polymerase III. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2013, 1829, 318-330.	1.9	98
16	Control of Transfer RNA Maturation by Phosphorylation of the Human La Antigen on Serine 366. <i>Molecular Cell</i> , 2000, 6, 339-348.	9.7	92
17	A conserved mechanism of TOR-dependent RCK-mediated mRNA degradation regulates autophagy. <i>Nature Cell Biology</i> , 2015, 17, 930-942.	10.3	91
18	La-Related Protein 4 Binds Poly(A), Interacts with the Poly(A)-Binding Protein MLE Domain via a Variant PAM2w Motif, and Can Promote mRNA Stability. <i>Molecular and Cellular Biology</i> , 2011, 31, 542-556.	2.3	89

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19	RNA Polymerase III Output Is Functionally Linked to tRNA Dimethyl-G26 Modification. <i>PLoS Genetics</i> , 2015, 11, e1005671.	3.5	81
20	Mechanism of Transcription Termination by RNA Polymerase III Utilizes a Non-template Strand Sequence-Specific Signal Element. <i>Molecular Cell</i> , 2015, 58, 1124-1132.	9.7	80
21	RNA Polymerase III Advances: Structural and tRNA Functional Views. <i>Trends in Biochemical Sciences</i> , 2016, 41, 546-559.	7.5	78
22	A novel perivascular cell population in the zebrafish brain. <i>ELife</i> , 2017, 6, .	6.0	77
23	Separate RNA-binding surfaces on the multifunctional La protein mediate distinguishable activities in tRNA maturation. <i>Nature Structural and Molecular Biology</i> , 2006, 13, 611-618.	8.2	76
24	The subset of mouse B1 (Alu-equivalent) sequences expressed as small processed cytoplasmic transcripts. <i>Nucleic Acids Research</i> , 1991, 19, 5695-5702.	14.5	74
25	Transcription Termination by RNA Polymerase III in Fission Yeast. <i>Journal of Biological Chemistry</i> , 2000, 275, 29076-29081.	3.4	71
26	Lack of tRNA Modification Isopentenyl-A37 Alters mRNA Decoding and Causes Metabolic Deficiencies in Fission Yeast. <i>Molecular and Cellular Biology</i> , 2013, 33, 2918-2929.	2.3	65
27	Comparative overview of RNA polymerase II and III transcription cycles, with focus on RNA polymerase III termination and reinitiation. <i>Transcription</i> , 2014, 5, e27369.	3.1	65
28	Evolving specificity of tRNA 3-methyl-cytidine-32 (m <sup>3</sup> C32) modification: a subset of tRNAs <sup>Ser</sup> requires N <sup>6</sup> -isopentenylation of A37. <i>Rna</i> , 2016, 22, 1400-1410.	3.5	64
29	Aberrant Nuclear Trafficking of La Protein Leads to Disordered Processing of Associated Precursor tRNAs. <i>Molecular Cell</i> , 2002, 9, 1113-1123.	9.7	63
30	Mutation of RNA Pol III Subunit <i>rpc2/polr3b</i> Leads to Deficiency of Subunit <i>Rpc11</i> and Disrupts Zebrafish Digestive Development. <i>PLoS Biology</i> , 2007, 5, e312.	5.6	63
31	Widespread Use of TATA Elements in the Core Promoters for RNA Polymerases III, II, and I in Fission Yeast. <i>Molecular and Cellular Biology</i> , 2001, 21, 6870-6881.	2.3	62
32	CK2 Is Responsible for Phosphorylation of Human La Protein Serine-366 and Can Modulate rpl37 5'-Terminal Oligopyrimidine mRNA Metabolism. <i>Molecular and Cellular Biology</i> , 2004, 24, 9580-9591.	2.3	62
33	Precursor-product discrimination by La protein during tRNA metabolism. <i>Nature Structural and Molecular Biology</i> , 2009, 16, 430-437.	8.2	57
34	Mutations in the RNA Polymerase III Subunit <i>Rpc11p</i> That Decrease RNA 5'-Terminal Oligo(U) Length and La-Dependent tRNA Processing. <i>Molecular and Cellular Biology</i> , 2005, 25, 621-636.	2.3	56
35	Global "bootprinting" reveals the elastic architecture of the yeast TFIIB-TFIIC transcription complex in vivo. <i>Nucleic Acids Research</i> , 2013, 41, 8135-8143.	14.5	55
36	The Multifunctional RNA-Binding Protein La Is Required for Mouse Development and for the Establishment of Embryonic Stem Cells. <i>Molecular and Cellular Biology</i> , 2006, 26, 1445-1451.	2.3	53

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37	Cellular La Protein Shields Nonsegmented Negative-Strand RNA Viral Leader RNA from RIG-I and Enhances Virus Growth by Diverse Mechanisms. <i>Journal of Virology</i> , 2008, 82, 7977-7987.	3.4	53
38	tRNA gene copy number variation in humans. <i>Gene</i> , 2014, 536, 376-384.	2.2	52
39	Monomeric scAlu and nascent dimeric Alu RNAs induced by adenovirus are assembled into SRP9/14-containing RNPs in HeLa cells. <i>Nucleic Acids Research</i> , 1996, 24, 4165-4170.	14.5	51
40	The La Protein-RNA Complex Surfaces. <i>Molecular Cell</i> , 2006, 21, 149-152.	9.7	49
41	Human Signal Recognition Particle (SRP) Alu-associated Protein Also Binds Alu Interspersed Repeat Sequence RNAs. <i>Journal of Biological Chemistry</i> , 1995, 270, 10179-10186.	3.4	47
42	Human La is found at RNA polymerase III-transcribed genes in vivo. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 18350-18355.	7.1	47
43	tRNAomics: tRNA gene copy number variation and codon use provide bioinformatic evidence of a new anticodon:codon wobble pair in a eukaryote. <i>Rna</i> , 2012, 18, 1358-1372.	3.5	47
44	The human Y4 small cytoplasmic RNA gene is controlled by upstream elements and resides on chromosome 7 with all other hY scRNA genes. <i>Nucleic Acids Research</i> , 1994, 22, 3045-3052.	14.5	43
45	Heterodimer SRP9/14 is an integral part of the neural BC200 RNP in primate brain. <i>Neuroscience Letters</i> , 1998, 245, 123-126.	2.1	43
46	LARP4 mRNA codon-tRNA match contributes to LARP4 activity for ribosomal protein mRNA poly(A) tail length protection. <i>ELife</i> , 2017, 6, .	6.0	43
47	La protein and its associated small nuclear and nucleolar precursor RNAs. <i>Gene Expression</i> , 2002, 10, 41-57.	1.2	43
48	Human Cells Have a Limited Set of tRNA Anticodon Loop Substrates of the tRNA Isopentenyltransferase TRIT1 Tumor Suppressor. <i>Molecular and Cellular Biology</i> , 2013, 33, 4900-4908.	2.3	42
49	Plasticity and diversity of tRNA anticodon determinants of substrate recognition by eukaryotic A37 isopentenyltransferases. <i>Rna</i> , 2011, 17, 1846-1857.	3.5	40
50	La Protein and the Trafficking of Nascent RNA Polymerase III Transcripts. <i>Journal of Cell Biology</i> , 2001, 153, F13-F18.	5.2	38
51	A chromosome 17q <i>de novo</i> paracentric inversion in a patient with campomelic dysplasia; case report and etiologic hypothesis. <i>Clinical Genetics</i> , 1991, 39, 401-408.	2.0	38
52	RNA polymerase III mutants in TFIIIF-like C37 that cause terminator readthrough with no decrease in transcription output. <i>Nucleic Acids Research</i> , 2013, 41, 139-155.	14.5	38
53	Nonphosphorylated Human La Antigen Interacts with Nucleolin at Nucleolar Sites Involved in rRNA Biogenesis. <i>Molecular and Cellular Biology</i> , 2004, 24, 10894-10904.	2.3	37
54	Point mutations in the Rpb9-homologous domain of Rpc11 that impair transcription termination by RNA polymerase III. <i>Nucleic Acids Research</i> , 2011, 39, 6100-6113.	14.5	35

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55	A call for direct sequencing of full-length RNAs to identify all modifications. <i>Nature Genetics</i> , 2021, 53, 1113-1116.	21.4	33
56	Conservation of a Masked Nuclear Export Activity of La Proteins and Its Effects on tRNA Maturation. <i>Molecular and Cellular Biology</i> , 2007, 27, 3303-3312.	2.3	31
57	Lack of tRNA-i6A modification causes mitochondrial-like metabolic deficiency in <i>S. pombe</i> by limiting activity of cytosolic tRNA <sup>Tyr</sup> , not mito-tRNA. <i>Rna</i> , 2016, 22, 583-596.	3.5	30
58	Factors That Shape Eukaryotic tRNAomes: Processing, Modification and Anticodon Use. <i>Biomolecules</i> , 2017, 7, 26.	4.0	30
59	Terminator-specific Recycling of a B1-Alu Transcription Complex by RNA Polymerase III Is Mediated by the RNA Terminus-binding Protein La. <i>Journal of Biological Chemistry</i> , 1998, 273, 26110-26116.	3.4	28
60	Different types of secondary information in the genetic code. <i>Rna</i> , 2014, 20, 977-984.	3.5	28
61	Distinguishing Core and Holoenzyme Mechanisms of Transcription Termination by RNA Polymerase III. <i>Molecular and Cellular Biology</i> , 2013, 33, 1571-1581.	2.3	27
62	Isolation and Cloning of Four Subunits of a Fission Yeast TFIIC Complex That Includes an Ortholog of the Human Regulatory Protein TFIIC <sup>2</sup> . <i>Journal of Biological Chemistry</i> , 2000, 275, 31480-31487.	3.4	26
63	A highly conserved nucleotide in the Alu domain of SRP RNA mediates translation arrest through high affinity binding to SRP9/14. <i>Nucleic Acids Research</i> , 1997, 25, 1117-1122.	14.5	25
64	Targeted Deletion of the Gene Encoding the La Autoantigen (Sjögren's Syndrome Antigen B) in B Cells or the Frontal Brain Causes Extensive Tissue Loss. <i>Molecular and Cellular Biology</i> , 2014, 34, 123-131.	2.3	24
65	La involvement in tRNA and other RNA processing events including differences among yeast and other eukaryotes. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2018, 1861, 361-372.	1.9	24
66	RNA polymerase III subunits C37/53 modulate rU:dA hybrid 3' end dynamics during transcription termination. <i>Nucleic Acids Research</i> , 2019, 47, 310-327.	14.5	24
67	Single molecule poly(A) tail-seq shows LARP4 opposes deadenylation throughout mRNA lifespan with most impact on short tails. <i>ELife</i> , 2020, 9, .	6.0	24
68	Altered nuclear tRNA metabolism in La-deleted <i>Schizosaccharomyces pombe</i> is accompanied by a nutritional stress response involving Atf1p and Pcr1p that is suppressible by Xpo-t/Los1p. <i>Molecular Biology of the Cell</i> , 2012, 23, 480-491.	2.1	21
69	The nuclear and cytoplasmic activities of RNA polymerase III, and an evolving transcriptome for surveillance. <i>Nucleic Acids Research</i> , 2021, 49, 12017-12034.	14.5	21
70	Comparative whole genome sequencing reveals phenotypic tRNA gene duplication in spontaneous <i>Schizosaccharomyces pombe</i> La mutants. <i>Nucleic Acids Research</i> , 2011, 39, 4728-4742.	14.5	20
71	Methylphosphate Cap Structure in Small RNAs Reduces the Affinity of RNAs to La Protein. <i>Gene Expression</i> , 2002, 10, 243-253.	1.2	20
72	Comment on "Mechanism of eukaryotic RNA polymerase III transcription termination". <i>Science</i> , 2014, 345, 524-524.	12.6	19

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73	The isolated La-module of LARP1 mediates 3' poly(A) protection and mRNA stabilization, dependent on its intrinsic PAM2 binding to PABPC1. <i>RNA Biology</i> , 2021, 18, 275-289.	3.1	18
74	A methods review on use of nonsense suppression to study 3' end formation and other aspects of tRNA biogenesis. <i>Gene</i> , 2015, 556, 35-50.	2.2	17
75	LARP1 and LARP4: up close with PABP for mRNA 3' poly(A) protection and stabilization. <i>RNA Biology</i> , 2021, 18, 259-274.	3.1	16
76	Active Center Control of Termination by RNA Polymerase III and tRNA Gene Transcription Levels In Vivo. <i>PLoS Genetics</i> , 2016, 12, e1006253.	3.5	15
77	Mouse and Human La Proteins Differ in Kinase Substrate Activity and Activation Mechanism for tRNA Processing. <i>Gene Expression</i> , 2007, 14, 71-81.	1.2	13
78	A transcriptional specialist resolved. <i>Nature</i> , 2015, 528, 204-205.	27.8	13
79	Mechanism of RNA polymerase III termination-associated reinitiation-recycling conferred by the essential function of the N terminal-and-linker domain of the C11 subunit. <i>Nature Communications</i> , 2021, 12, 5900.	12.8	13
80	It's a mod mod tRNA world. <i>Nature Chemical Biology</i> , 2008, 4, 162-164.	8.0	12
81	Targeting mitochondrial and cytosolic substrates of TRIT1 isopentenyltransferase: Specificity determinants and tRNA-i6A37 profiles. <i>PLoS Genetics</i> , 2020, 16, e1008330.	3.5	12
82	Construction of FLAG and histidine tagging vectors for <i>Schizosaccharomyces pombe</i> . <i>Yeast</i> , 2001, 18, 463-468.	1.7	11
83	TOR-dependent post-transcriptional regulation of autophagy. <i>Autophagy</i> , 2015, 11, 2390-2392.	9.1	11
84	LARP4 Is Regulated by Tumor Necrosis Factor Alpha in a Tristetraprolin-Dependent Manner. <i>Molecular and Cellular Biology</i> , 2016, 36, 574-584.	2.3	10
85	Transcriptome-wide stability analysis uncovers LARP4-mediated NF- $\kappa$ B1 mRNA stabilization during T cell activation. <i>Nucleic Acids Research</i> , 2020, 48, 8724-8739.	14.5	10
86	The fission yeast TFIIB-related factor limits RNA polymerase III to a TATA-dependent pathway of TBP recruitment. <i>Nucleic Acids Research</i> , 2003, 31, 2108-2116.	14.5	9
87	A high density of cis-information terminates RNA Polymerase III on a 2-rail track. <i>RNA Biology</i> , 2016, 13, 166-171.	3.1	9
88	La Deletion from Mouse Brain Alters Pre-tRNA Metabolism and Accumulation of Pre-5.8S rRNA, with Neuron Death and Reactive Astrocytosis. <i>Molecular and Cellular Biology</i> , 2017, 37, .	2.3	8
89	Biochemical Analysis of Transcription Termination by RNA Polymerase III from Yeast <i>Saccharomyces cerevisiae</i> . <i>Methods in Molecular Biology</i> , 2015, 1276, 185-198.	0.9	7
90	The conserved 7SK snRNA gene localizes to human chromosome 6 by homolog exclusion probing of somatic cell hybrid RNA. <i>Nucleic Acids Research</i> , 1994, 22, 722-725.	14.5	6

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91	A versatile tRNA modification-sensitive northern blot method with enhanced performance. <i>Rna</i> , 2022, 28, 418-432.	3.5	6
92	RNA Polymerase III from the Fission Yeast, <i>Schizosaccharomyces pombe</i> . <i>Methods in Enzymology</i> , 2003, 370, 165-173.	1.0	4
93	Human manganese superoxide dismutase is readily detectable by a copper blotting technique. <i>Biochemical Medicine and Metabolic Biology</i> , 1991, 46, 406-415.	0.7	3
94	Itâ€™s Snoâ€™ing on Pol III at nuclear pores. <i>Genome Biology</i> , 2013, 14, 137.	9.6	2
95	The leucine-NH <sub>4</sub> <sup>+</sup> uptake regulator Any1 limits growth as part of a general amino acid control response to loss of La protein by fission yeast. <i>PLoS ONE</i> , 2021, 16, e0253494.	2.5	2
96	Single-molecule polyadenylated tail sequencing (SM-PAT-Seq) to measure polyA tail lengths transcriptome-wide. <i>Methods in Enzymology</i> , 2021, 655, 119-137.	1.0	2