

Joshua T Mendell

List of Publications by Year in descending order

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Version: 2024-02-01

76
papers

34,708
citations

31976

53
h-index

74163

75
g-index

82
all docs

82
docs citations

82
times ranked

45993
citing authors

#	ARTICLE	IF	CITATIONS
1	StringTie enables improved reconstruction of a transcriptome from RNA-seq reads. <i>Nature Biotechnology</i> , 2015, 33, 290-295.	17.5	8,385
2	Functional Classification and Experimental Dissection of Long Noncoding RNAs. <i>Cell</i> , 2018, 172, 393-407.	28.9	2,657
3	c-Myc-regulated microRNAs modulate E2F1 expression. <i>Nature</i> , 2005, 435, 839-843.	27.8	2,618
4	Transactivation of miR-34a by p53 Broadly Influences Gene Expression and Promotes Apoptosis. <i>Molecular Cell</i> , 2007, 26, 745-752.	9.7	1,844
5	c-Myc suppression of miR-23a/b enhances mitochondrial glutaminase expression and glutamine metabolism. <i>Nature</i> , 2009, 458, 762-765.	27.8	1,801
6	Therapeutic microRNA Delivery Suppresses Tumorigenesis in a Murine Liver Cancer Model. <i>Cell</i> , 2009, 137, 1005-1017.	28.9	1,634
7	MicroRNAs in Stress Signaling and Human Disease. <i>Cell</i> , 2012, 148, 1172-1187.	28.9	1,471
8	Widespread microRNA repression by Myc contributes to tumorigenesis. <i>Nature Genetics</i> , 2008, 40, 43-50.	21.4	1,203
9	miRiad Roles for the miR-17-92 Cluster in Development and Disease. <i>Cell</i> , 2008, 133, 217-222.	28.9	1,012
10	Augmentation of tumor angiogenesis by a Myc-activated microRNA cluster. <i>Nature Genetics</i> , 2006, 38, 1060-1065.	21.4	1,000
11	MicroRNA-126 regulates endothelial expression of vascular cell adhesion molecule 1. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 1516-1521.	7.1	925
12	Nonsense surveillance regulates expression of diverse classes of mammalian transcripts and mutes genomic noise. <i>Nature Genetics</i> , 2004, 36, 1073-1078.	21.4	744
13	Noncoding RNA NORAD Regulates Genomic Stability by Sequestering PUMILIO Proteins. <i>Cell</i> , 2016, 164, 69-80.	28.9	723
14	Essential metabolic, anti-inflammatory, and anti-tumorigenic functions of miR-122 in liver. <i>Journal of Clinical Investigation</i> , 2012, 122, 2871-2883.	8.2	666
15	A Hexanucleotide Element Directs MicroRNA Nuclear Import. <i>Science</i> , 2007, 315, 97-100.	12.6	626
16	microRNAs in Vertebrate Physiology and Human Disease. <i>Annual Review of Genomics and Human Genetics</i> , 2007, 8, 215-239.	6.2	400
17	P53-induced microRNA-107 inhibits HIF-1 and tumor angiogenesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 6334-6339.	7.1	398
18	MicroRNAs: Critical Regulators of Development, Cellular Physiology and Malignancy. <i>Cell Cycle</i> , 2005, 4, 1179-1184.	2.6	388

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19	miR-21: An Androgen Receptor-Regulated MicroRNA that Promotes Hormone-Dependent and Hormone-Independent Prostate Cancer Growth. <i>Cancer Research</i> , 2009, 69, 7165-7169.	0.9	377
20	Lin-28B transactivation is necessary for Myc-mediated let-7 repression and proliferation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 3384-3389.	7.1	355
21	MicroRNA miR-155 is a biomarker of early pancreatic neoplasia. <i>Cancer Biology and Therapy</i> , 2009, 8, 340-346.	3.4	288
22	Restitution of Tumor Suppressor MicroRNAs Using a Systemic Nanovector Inhibits Pancreatic Cancer Growth in Mice. <i>Molecular Cancer Therapeutics</i> , 2011, 10, 1470-1480.	4.1	279
23	When the Message Goes Awry. <i>Cell</i> , 2001, 107, 411-414.	28.9	274
24	Repression of the miR-143/145 cluster by oncogenic Ras initiates a tumor-promoting feed-forward pathway. <i>Genes and Development</i> , 2010, 24, 2754-2759.	5.9	273
25	Separable Roles for rent1/hUpf1 in Altered Splicing and Decay of Nonsense Transcripts. <i>Science</i> , 2002, 298, 419-422.	12.6	246
26	An Argonaute phosphorylation cycle promotes microRNA-mediated silencing. <i>Nature</i> , 2017, 542, 197-202.	27.8	232
27	Epigenetic Silencing of MicroRNA miR-107 Regulates Cyclin-Dependent Kinase 6 Expression in Pancreatic Cancer. <i>Pancreatology</i> , 2009, 9, 293-301.	1.1	197
28	Somatic mutations in DROSHA and DICER1 impair microRNA biogenesis through distinct mechanisms in Wilms tumours. <i>Nature Communications</i> , 2014, 5, 4802.	12.8	192
29	Regulated expression of microRNAs in normal and polycythemia vera erythropoiesis. <i>Experimental Hematology</i> , 2007, 35, 1657-1667.	0.4	191
30	An Essential Mesenchymal Function for miR-143/145 in Intestinal Epithelial Regeneration. <i>Cell</i> , 2014, 157, 1104-1116.	28.9	188
31	Novel Upf2p Orthologues Suggest a Functional Link between Translation Initiation and Nonsense Surveillance Complexes. <i>Molecular and Cellular Biology</i> , 2000, 20, 8944-8957.	2.3	147
32	A ubiquitin ligase mediates target-directed microRNA decay independently of tailing and trimming. <i>Science</i> , 2020, 370, .	12.6	135
33	Targeting a Long Noncoding RNA in Breast Cancer. <i>New England Journal of Medicine</i> , 2016, 374, 2287-2289.	27.0	131
34	Myc: Maestro of MicroRNAs. <i>Genes and Cancer</i> , 2010, 1, 568-575.	1.9	123
35	NORAD-induced Pumilio phase separation is required for genome stability. <i>Nature</i> , 2021, 595, 303-308.	27.8	123
36	Cell-cell contact globally activates microRNA biogenesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 7016-7021.	7.1	122

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37	Identification of miR-21 targets in breast cancer cells using a quantitative proteomic approach. <i>Proteomics</i> , 2009, 9, 1374-1384.	2.2	113
38	A resource for analysis of microRNA expression and function in pancreatic ductal adenocarcinoma cells. <i>Cancer Biology and Therapy</i> , 2009, 8, 2013-2024.	3.4	108
39	Circular reasoning: microRNAs and cell-cycle control. <i>Trends in Biochemical Sciences</i> , 2008, 33, 474-481.	7.5	102
40	Antisense-Mediated Transcript Knockdown Triggers Premature Transcription Termination. <i>Molecular Cell</i> , 2020, 77, 1044-1054.e3.	9.7	100
41	Loss of CHD1 Promotes Heterogeneous Mechanisms of Resistance to AR-Targeted Therapy via Chromatin Dysregulation. <i>Cancer Cell</i> , 2020, 37, 584-598.e11.	16.8	96
42	Genome-wide annotation of microRNA primary transcript structures reveals novel regulatory mechanisms. <i>Genome Research</i> , 2015, 25, 1401-1409.	5.5	91
43	Role of pri-miRNA tertiary structure in miR-17~92 miRNA biogenesis. <i>RNA Biology</i> , 2011, 8, 1105-1114.	3.1	85
44	Last step in the path of LDL cholesterol from lysosome to plasma membrane to ER is governed by phosphatidylserine. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 18521-18529.	7.1	84
45	Analysis of regulatory network topology reveals functionally distinct classes of microRNAs. <i>Nucleic Acids Research</i> , 2008, 36, 6494-6503.	14.5	81
46	A novel source for miR-21 expression through the alternative polyadenylation of VMP1 gene transcripts. <i>Nucleic Acids Research</i> , 2012, 40, 6821-6833.	14.5	79
47	eIF5B drives integrated stress response-dependent translation of PD-L1 in lung cancer. <i>Nature Cancer</i> , 2020, 1, 533-545.	13.2	73
48	PUMILIO hyperactivity drives premature aging of Norad-deficient mice. <i>ELife</i> , 2019, 8, .	6.0	65
49	Transcriptional Regulation of miR-31 by Oncogenic KRAS Mediates Metastatic Phenotypes by Repressing RASA1. <i>Molecular Cancer Research</i> , 2016, 14, 267-277.	3.4	61
50	KIT signaling regulates MITF expression through miRNAs in normal and malignant mast cell proliferation. <i>Blood</i> , 2011, 117, 3629-3640.	1.4	60
51	Tumor suppression by miR-26 overrides potential oncogenic activity in intestinal tumorigenesis. <i>Genes and Development</i> , 2014, 28, 2585-2590.	5.9	59
52	Identifying targets of miR-143 using a SILAC-based proteomic approach. <i>Molecular BioSystems</i> , 2010, 6, 1873.	2.9	58
53	PUMILIO, but not RBMX, binding is required for regulation of genomic stability by noncoding RNA NORAD. <i>ELife</i> , 2019, 8, .	6.0	55
54	c-Myb oncoprotein is an essential target of the dleu2 tumor suppressor microRNA cluster. <i>Cancer Biology and Therapy</i> , 2008, 7, 1758-1764.	3.4	54

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55	Functional integration of microRNAs into oncogenic and tumor suppressor pathways. <i>Cell Cycle</i> , 2008, 7, 2493-2499.	2.6	53
56	Precise let-7 expression levels balance organ regeneration against tumor suppression. <i>ELife</i> , 2015, 4, e09431.	6.0	53
57	miR-26 suppresses adipocyte progenitor differentiation and fat production by targeting <i>Fbxl19</i> . <i>Genes and Development</i> , 2019, 33, 1367-1380.	5.9	50
58	Suppression of Ribosomal Pausing by eIF5A Is Necessary to Maintain the Fidelity of Start Codon Selection. <i>Cell Reports</i> , 2019, 29, 3134-3146.e6.	6.4	44
59	Noncoding RNAs and Cancer. <i>Cell</i> , 2013, 153, 9-10.	28.9	40
60	Mutations in microRNA processing genes in Wilms tumors derepress the <i>IGF2</i> regulator <i>PLAG1</i> . <i>Genes and Development</i> , 2018, 32, 996-1007.	5.9	40
61	Abate and Switch: miR-145 in Stem Cell Differentiation. <i>Cell</i> , 2009, 137, 606-608.	28.9	38
62	Systemic Delivery of scAAV8-Encoded MiR-29a Ameliorates Hepatic Fibrosis in Carbon Tetrachloride-Treated Mice. <i>PLoS ONE</i> , 2015, 10, e0124411.	2.5	37
63	Ribosome Recycling by ABCE1 Links Lysosomal Function and Iron Homeostasis to 3' UTR-Directed Regulation and Nonsense-Mediated Decay. <i>Cell Reports</i> , 2020, 32, 107895.	6.4	36
64	Loss of <i>Dis3l2</i> partially phenocopies Perlman syndrome in mice and results in up-regulation of <i>Igf2</i> in nephron progenitor cells. <i>Genes and Development</i> , 2018, 32, 903-908.	5.9	34
65	Quantification of Global MicroRNA Abundance by Selective Isotachophoresis. <i>Analytical Chemistry</i> , 2010, 82, 9631-9635.	6.5	31
66	MIR205HG Is a Long Noncoding RNA that Regulates Growth Hormone and Prolactin Production in the Anterior Pituitary. <i>Developmental Cell</i> , 2019, 49, 618-631.e5.	7.0	30
67	MicroRNA turnover: a tale of tailing, trimming, and targets. <i>Trends in Biochemical Sciences</i> , 2023, 48, 26-39.	7.5	28
68	Safety and Efficacy of AAV Retrograde Pancreatic Ductal Gene Delivery in Normal and Pancreatic Cancer Mice. <i>Molecular Therapy - Methods and Clinical Development</i> , 2018, 8, 8-20.	4.1	23
69	Identification of miR-145 targets through an integrated omics analysis. <i>Molecular BioSystems</i> , 2015, 11, 197-207.	2.9	21
70	Noncoding RNAs: biology and applications—a Keystone Symposia report. <i>Annals of the New York Academy of Sciences</i> , 2021, 1506, 118-141.	3.8	13
71	RBM33 directs the nuclear export of transcripts containing GC-rich elements. <i>Genes and Development</i> , 2022, 36, 550-565.	5.9	12
72	Tumors line up for a letdown. <i>Nature Genetics</i> , 2009, 41, 768-769.	21.4	11

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73	High-Throughput Characterization of Primary microRNA Transcripts. <i>Methods in Molecular Biology</i> , 2018, 1823, 1-9.	0.9	3
74	Seeing through the miRage of tissue complexity. <i>Cell Cycle</i> , 2014, 13, 2988-2989.	2.6	0
75	Dysregulated Expression of miRNAs in Polycythemia Vera Erythroid Progenitors. <i>Blood</i> , 2006, 108, 3613-3613.	1.4	0
76	Abstract P5-17-09: A genome-wide CRISPR screen identifies PRMT5 as a novel therapeutic target in ER+/ <i>RB1</i> -deficient breast cancer. <i>Cancer Research</i> , 2022, 82, P5-17-09-P5-17-09.	0.9	0