

# Yaron Shav-Tal

## List of Publications by Year in descending order

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Version: 2024-02-01

98  
papers

5,747  
citations

159585

30  
h-index

82547

72  
g-index

101  
all docs

101  
docs citations

101  
times ranked

6766  
citing authors

#	ARTICLE	IF	CITATIONS
1	From Silencing to Gene Expression. <i>Cell</i> , 2004, 116, 683-698.	28.9	658
2	In vivo dynamics of RNA polymerase II transcription. <i>Nature Structural and Molecular Biology</i> , 2007, 14, 796-806.	8.2	603
3	Dynamics of Single mRNPs in Nuclei of Living Cells. <i>Science</i> , 2004, 304, 1797-1800.	12.6	476
4	A transgenic mouse for in vivo detection of endogenous labeled mRNA. <i>Nature Methods</i> , 2011, 8, 165-170.	19.0	340
5	Dynamic Sorting of Nuclear Components into Distinct Nucleolar Caps during Transcriptional Inhibition. <i>Molecular Biology of the Cell</i> , 2005, 16, 2395-2413.	2.1	304
6	PSF and p54nrb/NonO - multi-functional nuclear proteins. <i>FEBS Letters</i> , 2002, 531, 109-114.	2.8	296
7	Dynamics of single mRNP nucleocytoplasmic transport and export through the nuclear pore in living cells. <i>Nature Cell Biology</i> , 2010, 12, 543-552.	10.3	230
8	The Dynamics of Mammalian P Body Transport, Assembly, and Disassembly In Vivo. <i>Molecular Biology of the Cell</i> , 2008, 19, 4154-4166.	2.1	208
9	The In Vivo Kinetics of RNA Polymerase II Elongation during Co-Transcriptional Splicing. <i>PLoS Biology</i> , 2011, 9, e1000573.	5.6	171
10	Stepwise RNP assembly at the site of H/ACA RNA transcription in human cells. <i>Journal of Cell Biology</i> , 2006, 173, 207-218.	5.2	161
11	Single-allele analysis of transcription kinetics in living mammalian cells. <i>Nature Methods</i> , 2010, 7, 631-633.	19.0	155
12	The life of an mRNA in space and time. <i>Journal of Cell Science</i> , 2010, 123, 1761-1774.	2.0	112
13	Quantifying mRNA targeting to P bodies in living human cells reveals a dual role in mRNA decay and storage. <i>Journal of Cell Science</i> , 2014, 127, 4443-56.	2.0	106
14	Imaging gene expression in single living cells. <i>Nature Reviews Molecular Cell Biology</i> , 2004, 5, 855-862.	37.0	105
15	The Dbp5 cycle at the nuclear pore complex during mRNA export I: <i>dbp5</i> mutants with defects in RNA binding and ATP hydrolysis define key steps for Nup159 and Gle1. <i>Genes and Development</i> , 2011, 25, 1052-1064.	5.9	99
16	The Plasmacytoma Growth Inhibitor Restrictin-P Is an Antagonist of Interleukin 6 and Interleukin 11. <i>Journal of Biological Chemistry</i> , 1995, 270, 29594-29600.	3.4	88
17	The differential interaction of snRNPs with pre-mRNA reveals splicing kinetics in living cells. <i>Journal of Cell Biology</i> , 2010, 191, 75-86.	5.2	87
18	Assembling an intermediate filament network by dynamic cotranslation. <i>Journal of Cell Biology</i> , 2006, 172, 747-758.	5.2	74

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19	RNA localization. <i>Journal of Cell Science</i> , 2005, 118, 4077-4081.	2.0	69
20	The Role of Activin A in Regulation of Hemopoiesis. <i>Stem Cells</i> , 2002, 20, 493-500.	3.2	66
21	The nuclear cap-binding complex interacts with the U4/U6-U5 tri-snRNP and promotes spliceosome assembly in mammalian cells. <i>Rna</i> , 2013, 19, 1054-1063.	3.5	65
22	Exploring chromatin organization mechanisms through its dynamic properties. <i>Nucleus</i> , 2016, 7, 27-33.	2.2	63
23	Quantifying $\beta$ -catenin subcellular dynamics and cyclin D1 mRNA transcription during Wnt signaling in single living cells. <i>ELife</i> , 2016, 5, .	6.0	58
24	Nuclear Relocalization of the Pre-mRNA Splicing Factor PSF during Apoptosis Involves Hyperphosphorylation, Masking of Antigenic Epitopes, and Changes in Protein Interactions. <i>Molecular Biology of the Cell</i> , 2001, 12, 2328-2340.	2.1	52
25	The P Body Protein Dcp1a Is Hyper-phosphorylated during Mitosis. <i>PLoS ONE</i> , 2013, 8, e49783.	2.5	52
26	Dynamics of transcription and mRNA export. <i>Current Opinion in Cell Biology</i> , 2005, 17, 332-339.	5.4	45
27	A hydrothermal reaction of an aqueous solution of BSA yields highly fluorescent N doped C-dots used for imaging of live mammalian cells. <i>Journal of Materials Chemistry B</i> , 2016, 4, 2913-2920.	5.8	45
28	Cellular Levels of Signaling Factors Are Sensed by $\beta$ -actin Alleles to Modulate Transcriptional Pulse Intensity. <i>Cell Reports</i> , 2015, 11, 419-432.	6.4	41
29	Uncoupling of nucleo-cytoplasmic RNA export and localization during stress. <i>Nucleic Acids Research</i> , 2019, 47, 4778-4797.	14.5	39
30	Adhesion molecules involved in the interactions between early T cells and mesenchymal bone marrow stromal cells. <i>Experimental Hematology</i> , 1999, 27, 834-844.	0.4	38
31	Sonochemical Synthesis of DNA Nanospheres. <i>ChemBioChem</i> , 2011, 12, 1678-1681.	2.6	32
32	Gene expression within a dynamic nuclear landscape. <i>EMBO Journal</i> , 2006, 25, 3469-3479.	7.8	30
33	Intracellular trafficking and dynamics of P bodies. <i>Prion</i> , 2008, 2, 131-134.	1.8	30
34	Availability of splicing factors in the nucleoplasm can regulate the release of mRNA from the gene after transcription. <i>PLoS Genetics</i> , 2019, 15, e1008459.	3.5	29
35	Into the basket and beyond: the journey of mRNA through the nuclear pore complex. <i>Biochemical Journal</i> , 2020, 477, 23-44.	3.7	29
36	Quantifying the transcriptional output of single alleles in single living mammalian cells. <i>Nature Protocols</i> , 2013, 8, 393-408.	12.0	27

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37	The proteolysis adaptor, NblA, is essential for degradation of the core pigment of the cyanobacterial light-harvesting complex. <i>Plant Journal</i> , 2015, 83, 845-852.	5.7	27
38	Gene architecture directs splicing outcome in separate nuclear spatial regions. <i>Molecular Cell</i> , 2022, 82, 1021-1034.e8.	9.7	26
39	mRNPs meet stress granules. <i>FEBS Letters</i> , 2017, 591, 2534-2542.	2.8	25
40	Imaging within single NPCs reveals NXF1's role in mRNA export on the cytoplasmic side of the pore. <i>Journal of Cell Biology</i> , 2019, 218, 2962-2981.	5.2	24
41	Nuclear speckles – a driving force in gene expression. <i>Journal of Cell Science</i> , 2022, 135, .	2.0	23
42	The proteolysis adaptor, NblA, initiates protein pigment degradation by interacting with the cyanobacterial light-harvesting complexes. <i>Plant Journal</i> , 2014, 79, 118-126.	5.7	22
43	De-novo protein function prediction using DNA binding and RNA binding proteins as a test case. <i>Nature Communications</i> , 2016, 7, 13424.	12.8	22
44	Reorganization of nuclear factors during myeloid differentiation. <i>Journal of Cellular Biochemistry</i> , 2001, 81, 379-392.	2.6	20
45	The mesenchyme expresses T cell receptor mRNAs: relevance to cell growth control. <i>Oncogene</i> , 2002, 21, 2029-2036.	5.9	20
46	The Mesenchymal Stroma Negatively Regulates B Cell Lymphopoiesis through the Expression of Activin A. <i>Annals of the New York Academy of Sciences</i> , 2003, 996, 245-260.	3.8	20
47	Resolving the spatial relationship between intracellular components by dual color super resolution optical fluctuations imaging (SOFI). <i>Optical Nanoscopy</i> , 2013, 2, .	4.0	20
48	Enhanced proteolysis of pre-mRNA splicing factors in myeloid cells. <i>Experimental Hematology</i> , 2000, 28, 1029-1038.	0.4	18
49	Transcription and splicing. <i>Transcription</i> , 2011, 2, 216-220.	3.1	18
50	Peroxisome function relies on organelle-associated mRNA translation. <i>Science Advances</i> , 2022, 8, eabk2141.	10.3	18
51	Binding properties and dynamic localization of an alternative isoform of the cap-binding complex subunit CBP20. <i>Nucleus</i> , 2010, 1, 412-421.	2.2	17
52	Mutations in S-adenosylhomocysteine hydrolase (AHCY) affect its nucleocytoplasmic distribution and capability to interact with S-adenosylhomocysteine hydrolase-like 1 protein. <i>European Journal of Cell Biology</i> , 2017, 96, 579-590.	3.6	17
53	Proteinaceous microspheres for targeted RNA delivery prepared by an ultrasonic emulsification method. <i>Journal of Materials Chemistry B</i> , 2013, 1, 82-90.	5.8	16
54	Dynamic Encounters of Genes and Transcripts with the Nuclear Pore. <i>Trends in Genetics</i> , 2016, 32, 419-431.	6.7	16

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55	Applying styryl quinolinium fluorescent probes for imaging of ribosomal RNA in living cells. <i>Dyes and Pigments</i> , 2020, 174, 107986.	3.7	16
56	Dynamics and kinetics of nucleocytoplasmic mRNA export. <i>Wiley Interdisciplinary Reviews RNA</i> , 2010, 1, 388-401.	6.4	15
57	Speculating on the Roles of Nuclear Speckles: How RNA-Protein Nuclear Assemblies Affect Gene Expression. <i>BioEssays</i> , 2020, 42, e2000104.	2.5	15
58	Detection of mRNA of the Cyclin D1 Breast Cancer Marker by a Novel Duplex-DNA Probe. <i>Journal of Medicinal Chemistry</i> , 2013, 56, 4860-4869.	6.4	14
59	The dynamic pathway of nuclear RNA in eukaryotes. <i>Nucleus</i> , 2013, 4, 195-205.	2.2	13
60	Phospho-Tau Impairs Nuclear-Cytoplasmic Transport. <i>ACS Chemical Neuroscience</i> , 2019, 10, 36-38.	3.5	12
61	CD-tagging-MS2: detecting allelic expression of endogenous mRNAs and their protein products in single cells. <i>Biology Methods and Protocols</i> , 2017, 2, bpx004.	2.2	11
62	The dynamic lifecycle of mRNA in the nucleus. <i>Current Opinion in Cell Biology</i> , 2019, 58, 69-75.	5.4	11
63	Single-molecule dynamics of nuclear mRNA. <i>F1000 Biology Reports</i> , 2009, 1, 29.	4.0	11
64	Incomplete T-cell receptor <sup>12</sup> peptides target the mitochondrion and induce apoptosis. <i>Blood</i> , 2009, 113, 3530-3541.	1.4	10
65	Visualizing nuclear RNAi activity in single living human cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E8837-E8846.	7.1	10
66	An oligonucleotide probe incorporating the chromophore of green fluorescent protein is useful for the detection of HER-2 mRNA breast cancer marker. <i>European Journal of Medicinal Chemistry</i> , 2019, 173, 99-106.	5.5	10
67	The Portal Vertex of KSHV Promotes Docking of Capsids at the Nuclear Pores. <i>Viruses</i> , 2021, 13, 597.	3.3	10
68	The living test-tube: imaging of real-time gene expression. <i>Soft Matter</i> , 2006, 2, 361.	2.7	9
69	Detection of cyclin D1 mRNA by hybridization sensitive NIC oligonucleotide probe. <i>Bioorganic and Medicinal Chemistry</i> , 2014, 22, 2613-2621.	3.0	9
70	Cytoplasmic DNA can be detected by RNA fluorescence in situ hybridization. <i>Nucleic Acids Research</i> , 2019, 47, e109-e109.	14.5	9
71	Glucocorticoids enhance chemotherapy-driven stress granule assembly and impair granule dynamics, leading to cell death. <i>Journal of Cell Science</i> , 2022, 135, .	2.0	9
72	Single-site transcription rates through fitting of ensemble-averaged data from fluorescence recovery after photobleaching: A fat-tailed distribution. <i>Physical Review E</i> , 2015, 92, 032715.	2.1	8

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73	On the right track. <i>Nucleus</i> , 2010, 1, 492-498.	2.2	7
74	FRET energy transfer via Pdots improves the efficiency of photodynamic therapy and leads to rapid cell death. <i>Journal of Photochemistry and Photobiology B: Biology</i> , 2016, 164, 123-131.	3.8	7
75	Specific, Sensitive, and Quantitative Detection of HER-2 mRNA Breast Cancer Marker by Fluorescent Light-Up Hybridization Probes. <i>Bioconjugate Chemistry</i> , 2020, 31, 1188-1198.	3.6	7
76	Zooming in on single active genes in living mammalian cells. <i>Histochemistry and Cell Biology</i> , 2013, 140, 71-79.	1.7	6
77	Acting on impulse: dissecting the dynamics of the NFAT transcriptional response. <i>Genome Biology</i> , 2013, 14, 102.	9.6	6
78	The dynamics of the alternatively spliced NOL7 gene products and role in nucleolar architecture. <i>Nucleus</i> , 2011, 2, 229-245.	2.2	5
79	Development of fluorescent double-strand probes labeled with 8-(p-CF3-cinnamyl)-adenosine for the detection of cyclin D1 breast cancer marker. <i>European Journal of Medicinal Chemistry</i> , 2014, 79, 77-88.	5.5	5
80	The stress-inducible transcription factor ATF4 accumulates at specific rRNA-processing nucleolar regions after proteasome inhibition. <i>European Journal of Cell Biology</i> , 2016, 95, 389-400.	3.6	5
81	Yeast and Human Nuclear Pore Complexes: Not So Similar After All. <i>Trends in Cell Biology</i> , 2018, 28, 589-591.	7.9	5
82	S-phase transcriptional buffering quantified on two different promoters. <i>Life Science Alliance</i> , 2018, 1, e201800086.	2.8	5
83	Measuring the Kinetics of mRNA Transcription in Single Living Cells. <i>Journal of Visualized Experiments</i> , 2011, , e2898.	0.3	4
84	Single mRNP Tracking in Living Mammalian Cells. <i>Methods in Molecular Biology</i> , 2013, 1042, 87-99.	0.9	4
85	Measuring transcription dynamics in living cells using a photobleaching approach. <i>Methods</i> , 2017, 120, 58-64.	3.8	4
86	The Sub-Nuclear Localization of RNA-Binding Proteins in KSHV-Infected Cells. <i>Cells</i> , 2020, 9, 1958.	4.1	3
87	Dynamic Supraspliceosomes Are Assembled on Different Transcripts Regardless of Their Intron Number and Splicing State. <i>Frontiers in Genetics</i> , 2020, 11, 409.	2.3	3
88	Imaging mRNAs in Living Mammalian Cells. <i>Methods in Molecular Biology</i> , 2011, 714, 249-263.	0.9	3
89	Quantifying the Ratio of Spliceosome Components Assembled on Pre-mRNA. <i>Methods in Molecular Biology</i> , 2014, 1126, 257-269.	0.9	3
90	The Association of MEG3 lncRNA with Nuclear Speckles in Living Cells. <i>Cells</i> , 2022, 11, 1942.	4.1	3

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91	Visualizing transcription in real-time. <i>Open Life Sciences</i> , 2008, 3, 11-18.	1.4	2
92	Methods for visualizing RNA in cells, tissues and whole organisms. <i>Methods</i> , 2016, 98, 1-3.	3.8	2
93	Active RNA polymerase II curbs chromatin movement. <i>Journal of Cell Biology</i> , 2019, 218, 1427-1428.	5.2	2
94	Dynamics and Transport of Nuclear RNA. , 2016, , 491-513.		1
95	Detection of mRNAs Anchored to the Nuclear Envelope During Export Inhibition in Living Cells. <i>Methods in Molecular Biology</i> , 2019, 2038, 151-163.	0.9	1
96	Nuclear biology: making sense of complex processes. <i>Molecular Biology of the Cell</i> , 2012, 23, 976-976.	2.1	0
97	Visualizing Nuclear RNA Editing. <i>Trends in Biochemical Sciences</i> , 2017, 42, 845-847.	7.5	0
98	Dissecting Cellular Activity from Single Genes to Single mRNAs. , 0, , 29-39.		0