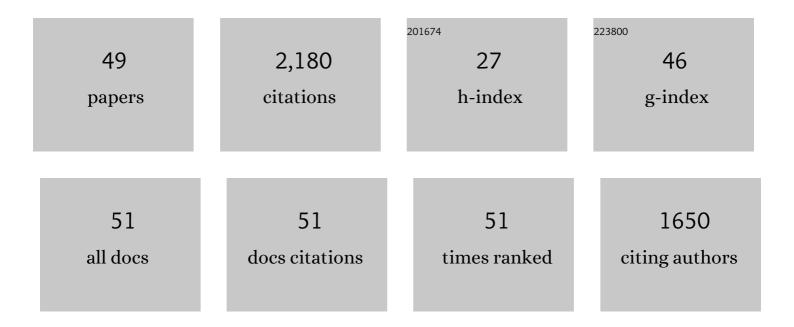
## Alice Telesnitsky

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Bimodal Expression Patterns, and Not Viral Burst Sizes, Predict the Effects of Vpr on HIV-1 Proviral Populations in Jurkat Cells. MBio, 2022, , e0374821.	4.1	0
2	Stability and conformation of the dimeric HIV-1 genomic RNA 5′UTR. Biophysical Journal, 2021, 120, 4874-4890.	0.5	13
3	5a€ <sup>2</sup> -Cap sequestration is an essential determinant of HIV-1 genome packaging. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	26
4	HIV-1 spliced RNAs display transcription start site bias. Rna, 2020, 26, 708-714.	3.5	11
5	Identification of the initial nucleocapsid recognition element in the HIV-1 RNA packaging signal. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 17737-17746.	7.1	50
6	Structural basis for transcriptional start site control of HIV-1 RNA fate. Science, 2020, 368, 413-417.	12.6	76
7	Flexibility in Nucleic Acid Binding Is Central to APOBEC3H Antiviral Activity. Journal of Virology, 2019, 93, .	3.4	8
8	Stable integrant-specific differences in bimodal HIV-1 expression patterns revealed by high-throughput analysis. PLoS Pathogens, 2019, 15, e1007903.	4.7	5
9	Influence of gag and RRE Sequences on HIV-1 RNA Packaging Signal Structure and Function. Journal of Molecular Biology, 2018, 430, 2066-2079.	4.2	21
10	Multiple, Switchable Protein:RNA Interactions Regulate Human Immunodeficiency Virus Type 1 Assembly. Annual Review of Virology, 2018, 5, 165-183.	6.7	50
11	The Host RNAs in Retroviral Particles. Viruses, 2016, 8, 235.	3.3	40
12	Host RNA Packaging by Retroviruses: A Newly Synthesized Story. MBio, 2016, 7, e02025-15.	4.1	32
13	Resolution of Specific Nucleotide Mismatches by Wild-Type and AZT-Resistant Reverse Transcriptases during HIV-1 Replication. Journal of Molecular Biology, 2016, 428, 2275-2288.	4.2	7
14	Transcriptional start site heterogeneity modulates the structure and function of the HIV-1 genome. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 13378-13383.	7.1	78
15	Analysis of the human immunodeficiency virus-1 RNA packageome. Rna, 2016, 22, 1228-1238.	3.5	46
16	Structure of the HIV-1 RNA packaging signal. Science, 2015, 348, 917-921.	12.6	211
17	A retrovirus packages nascent host noncoding RNAs from a novel surveillance pathway. Genes and Development, 2015, 29, 646-657.	5.9	40
18	Determinants of Moloney Murine Leukemia Virus Gag-Pol and Genomic RNA Proportions. Journal of Virology, 2014, 88, 7267-7275.	3.4	3

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#	Article	IF	CITATIONS
19	cis-Acting Determinants of 7SL RNA Packaging by HIV-1. Journal of Virology, 2012, 86, 7934-7942.	3.4	19
20	Identification of a Minimal Region of the HIV-1 5′-Leader Required for RNA Dimerization, NC Binding, and Packaging. Journal of Molecular Biology, 2012, 417, 224-239.	4.2	89
21	Moloney murine leukemia virus genomic RNA packaged in the absence of a full complement of wild type nucleocapsid protein. Virology, 2012, 430, 100-109.	2.4	3
22	NMR Detection of Structures in the HIV-1 5′-Leader RNA That Regulate Genome Packaging. Science, 2011, 334, 242-245.	12.6	227
23	7SL RNA Is Retained in HIV-1 Minimal Virus-Like Particles as an S-Domain Fragment. Journal of Virology, 2010, 84, 9070-9077.	3.4	34
24	Retroviral RNA Dimerization and Packaging: The What, How, When, Where, and Why. PLoS Pathogens, 2010, 6, e1001007.	4.7	83
25	Retroviruses: Molecular Biology, Genomics and Pathogenesis. Future Virology, 2010, 5, 539-543.	1.8	15
26	An RNA Structural Switch Regulates Diploid Genome Packaging by Moloney Murine Leukemia Virus. Journal of Molecular Biology, 2010, 396, 141-152.	4.2	46
27	Structure of a Conserved Retroviral RNA Packaging Element by NMR Spectroscopy and Cryo-Electron Tomography. Journal of Molecular Biology, 2010, 404, 751-772.	4.2	63
28	Packaging of Host mY RNAs by Murine Leukemia Virus May Occur Early in Y RNA Biogenesis. Journal of Virology, 2009, 83, 12526-12534.	3.4	37
29	The Remarkable Frequency of Human Immunodeficiency Virus Type 1 Genetic Recombination. Microbiology and Molecular Biology Reviews, 2009, 73, 451-480.	6.6	139
30	Pseudodiploid Genome Organization Aids Full-Length Human Immunodeficiency Virus Type 1 DNA Synthesis. Journal of Virology, 2008, 82, 2376-2384.	3.4	18
31	Effects of Identity Minimization on Moloney Murine Leukemia Virus Template Recognition and Frequent Tertiary Template-Directed Insertions during Nonhomologous Recombination. Journal of Virology, 2007, 81, 12156-12168.	3.4	5
32	Characterization of a natural heterodimer between MLV genomic RNA and the SDâ $\in^2$ retroelement generated by alternative splicing. Rna, 2007, 13, 2266-2276.	3.5	12
33	Two distinct Moloney murine leukemia virus RNAs produced from a single locus dimerize at random. Virology, 2006, 344, 391-400.	2.4	29
34	Evidence for the acquisition of multi-drug resistance in an HIV-1 clinical isolate via human sequence transduction. Virology, 2006, 351, 1-6.	2.4	9
35	7SL RNA, but not the 54-kd signal recognition particle protein, is an abundant component of both infectious HIV-1 and minimal virus-like particles. Rna, 2006, 12, 542-546.	3.5	85
36	Nonrandom Packaging of Host RNAs in Moloney Murine Leukemia Virus. Journal of Virology, 2005, 79, 13528-13537.	3.4	64

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#	Article	IF	CITATIONS
37	Nonrandom Dimerization of Murine Leukemia Virus Genomic RNAs. Journal of Virology, 2004, 78, 12129-12139.	3.4	41
38	Human Immunodeficiency Virus Type 1 Transductive Recombination Can Occur Frequently and in Proportion to Polyadenylation Signal Readthrough. Journal of Virology, 2004, 78, 3419-3428.	3.4	22
39	Mismatch Extension During Strong Stop Strand Transfer and Minimal Homology Requirements for Replicative Template Switching During Moloney Murine Leukemia Virus Replication. Journal of Molecular Biology, 2003, 330, 657-674.	4.2	8
40	Human Immunodeficiency Virus Type 1 Genetic Recombination Is More Frequent Than That of Moloney Murine Leukemia Virus despite Similar Template Switching Rates. Journal of Virology, 2003, 77, 4577-4587.	3.4	69
41	RNase H Activity Is Required for High-Frequency Repeat Deletion during Moloney Murine Leukemia Virus Replication. Journal of Virology, 2002, 76, 88-95.	3.4	26
42	Frequency of Direct Repeat Deletion in a Human Immunodeficiency Virus Type 1 Vector during Reverse Transcription in Human Cells. Virology, 2001, 286, 475-482.	2.4	43
43	Effects of Limiting Homology at the Site of Intermolecular Recombinogenic Template Switching during Moloney Murine Leukemia Virus Replication. Journal of Virology, 2001, 75, 11263-11274.	3.4	40
44	Structure-Based Moloney Murine Leukemia Virus Reverse Transcriptase Mutants with Altered Intracellular Direct-Repeat Deletion Frequencies. Journal of Virology, 2000, 74, 9629-9636.	3.4	22
45	Replication of Lengthened Moloney Murine Leukemia Virus Genomes Is Impaired at Multiple Stages. Journal of Virology, 2000, 74, 2694-2702.	3.4	37
46	Altering the Intracellular Environment Increases the Frequency of Tandem Repeat Deletion during Moloney Murine Leukemia Virus Reverse Transcription. Journal of Virology, 1999, 73, 8441-8447.	3.4	56
47	Effects of 3′ Untranslated Region Mutations on Plus-Strand Priming during Moloney Murine Leukemia Virus Replication. Journal of Virology, 1999, 73, 948-957.	3.4	27
48	Cis-acting elements required for strong stop acceptor template selection during moloney murine leukemia virus reverse transcription. Journal of Molecular Biology, 1998, 281, 1-15.	4.2	27
49	[27] Assays for retroviral reverse transcriptase. Methods in Enzymology, 1995, 262, 347-362.	1.0	68