

Judith G Levin

List of Publications by Year in descending order

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2,922
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236925

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1608
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#	ARTICLE	IF	CITATIONS
1	Mechanistic differences between HIV-1 and SIV nucleocapsid proteins and cross-species HIV-1 genomic RNA recognition. <i>Retrovirology</i> , 2016, 13, 89.	2.0	13
2	Nuclear Magnetic Resonance Structure of the APOBEC3B Catalytic Domain: Structural Basis for Substrate Binding and DNA Deaminase Activity. <i>Biochemistry</i> , 2016, 55, 2944-2959.	2.5	55
3	Sequence and structural determinants of human APOBEC3H deaminase and anti-HIV-1 activities. <i>Retrovirology</i> , 2015, 12, 3.	2.0	32
4	Structural determinants of human APOBEC3A enzymatic and nucleic acid binding properties. <i>Nucleic Acids Research</i> , 2014, 42, 1095-1110.	14.5	68
5	Oligomerization transforms human APOBEC3G from an efficient enzyme to a slowly dissociating nucleic acid-binding protein. <i>Nature Chemistry</i> , 2014, 6, 28-33.	13.6	67
6	Selection of fully processed HIV-1 nucleocapsid protein is required for optimal nucleic acid chaperone activity in reverse transcription. <i>Virus Research</i> , 2014, 193, 52-64.	2.2	13
7	Zinc finger function of HIV-1 nucleocapsid protein is required for removal of 5' terminal genomic RNA fragments: A paradigm for RNA removal reactions in HIV-1 reverse transcription. <i>Virus Research</i> , 2013, 171, 346-355.	2.2	9
8	Obituary. <i>Virus Research</i> , 2013, 171, 356.	2.2	0
9	The interdomain linker region of HIV-1 capsid protein is a critical determinant of proper core assembly and stability. <i>Virology</i> , 2011, 421, 253-265.	2.4	51
10	Fundamental differences between the nucleic acid chaperone activities of HIV-1 nucleocapsid protein and Gag or Gag-derived proteins: Biological implications. <i>Virology</i> , 2010, 405, 556-567.	2.4	41
11	Role of HIV-1 nucleocapsid protein in HIV-1 reverse transcription. <i>RNA Biology</i> , 2010, 7, 754-774.	3.1	141
12	Fidelity of plus-strand priming requires the nucleic acid chaperone activity of HIV-1 nucleocapsid protein. <i>Nucleic Acids Research</i> , 2009, 37, 1755-1766.	14.5	22
13	HIV-1 Vif-mediated ubiquitination/degradation of APOBEC3G involves four critical lysine residues in its C-terminal domain. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 19539-19544.	7.1	53
14	Effects of nucleic acid local structure and magnesium ions on minus-strand transfer mediated by the nucleic acid chaperone activity of HIV-1 nucleocapsid protein. <i>Nucleic Acids Research</i> , 2007, 35, 3974-3987.	14.5	23
15	Deaminase-independent inhibition of HIV-1 reverse transcription by APOBEC3G. <i>Nucleic Acids Research</i> , 2007, 35, 7096-7108.	14.5	281
16	A second-site suppressor significantly improves the defective phenotype imposed by mutation of an aromatic residue in the N-terminal domain of the HIV-1 capsid protein. <i>Virology</i> , 2007, 359, 105-115.	2.4	10
17	Biochemical Activities of Highly Purified, Catalytically Active Human APOBEC3G: Correlation with Antiviral Effect. <i>Journal of Virology</i> , 2006, 80, 5992-6002.	3.4	184
18	Effect of Polypurine Tract (PPT) Mutations on Human Immunodeficiency Virus Type 1 Replication: a Virus with a Completely Randomized PPT Retains Low Infectivity. <i>Journal of Virology</i> , 2005, 79, 6859-6867.	3.4	23

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19	Nucleic Acid Chaperone Activity of HIV-1 Nucleocapsid Protein: Critical Role in Reverse Transcription and Molecular Mechanism. <i>Progress in Molecular Biology and Translational Science</i> , 2005, 80, 217-286.	1.9	302
20	Alteration of Nucleic Acid Structure and Stability Modulates the Efficiency of Minus-Strand Transfer Mediated by the HIV-1 Nucleocapsid Protein. <i>Journal of Biological Chemistry</i> , 2004, 279, 44154-44165.	3.4	48
21	Nucleic Acid Conformational Changes Essential for HIV-1 Nucleocapsid Protein-mediated Inhibition of Self-priming in Minus-strand Transfer. <i>Journal of Molecular Biology</i> , 2003, 325, 1-10.	4.2	67
22	Human Immunodeficiency Virus Type 1 N-Terminal Capsid Mutants Containing Cores with Abnormally High Levels of Capsid Protein and Virtually No Reverse Transcriptase. <i>Journal of Virology</i> , 2003, 77, 12592-12602.	3.4	50
23	Efficient Initiation of HIV-1 Reverse Transcription <i>In Vitro</i> . <i>Journal of Biological Chemistry</i> , 2003, 278, 14185-14195.	3.4	44
24	Subtle Alterations of the Native Zinc Finger Structures Have Dramatic Effects on the Nucleic Acid Chaperone Activity of Human Immunodeficiency Virus Type 1 Nucleocapsid Protein. <i>Journal of Virology</i> , 2002, 76, 4370-4378.	3.4	100
25	Human Immunodeficiency Virus Type 1 N-Terminal Capsid Mutants That Exhibit Aberrant Core Morphology and Are Blocked in Initiation of Reverse Transcription in Infected Cells. <i>Journal of Virology</i> , 2001, 75, 9357-9366.	3.4	135
26	Zinc Finger Structures in the Human Immunodeficiency Virus Type 1 Nucleocapsid Protein Facilitate Efficient Minus- and Plus-Strand Transfer. <i>Journal of Virology</i> , 2000, 74, 8980-8988.	3.4	192
27	A Mechanism for Plus-Strand Transfer Enhancement by the HIV-1 Nucleocapsid Protein during Reverse Transcription. <i>Biochemistry</i> , 2000, 39, 9084-9091.	2.5	94
28	Molecular Requirements for Human Immunodeficiency Virus Type 1 Plus-Strand Transfer: Analysis in Reconstituted and Endogenous Reverse Transcription Systems. <i>Journal of Virology</i> , 1999, 73, 4794-4805.	3.4	63
29	Nucleic-acid-chaperone activity of retroviral nucleocapsid proteins: significance for viral replication. <i>Trends in Biochemical Sciences</i> , 1998, 23, 297-301.	7.5	370
30	Generation of HIV-1/HIV-2 cross-reactive peptide antisera by small sequence changes in HIV-1 reverse transcriptase and integrase immunizing peptides. <i>Journal of Biomedical Science</i> , 1998, 5, 192-202.	7.0	8
31	Actinomycin D Inhibits Human Immunodeficiency Virus Type 1 Minus-Strand Transfer in <i>In Vitro</i> and Endogenous Reverse Transcriptase Assays. <i>Journal of Virology</i> , 1998, 72, 6716-6724.	3.4	68
32	Mutating a Conserved Motif of the HIV-1 Reverse Transcriptase Palm Subdomain Alters Primer Utilization. <i>Biochemistry</i> , 1997, 36, 5758-5768.	2.5	55
33	Defects in Primer-Template Binding, Processive DNA Synthesis, and RNase H Activity Associated with Chimeric Reverse Transcriptases Having the Murine Leukemia Virus Polymerase Domain Joined to <i>Escherichia coli</i> RNase H. <i>Biochemistry</i> , 1995, 34, 5018-5029.	2.5	54
34	Deficiency of 60 to 70 S RNA in Murine Leukemia Virus Particles Assembled in Cells Treated with Actinomycin D. <i>Journal of Virology</i> , 1974, 14, 152-161.	3.4	186