

# Andrew P Rice

## List of Publications by Year in descending order

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85  
papers

4,174  
citations

126907

33  
h-index

123424

61  
g-index

87  
all docs

87  
docs citations

87  
times ranked

3908  
citing authors

#	ARTICLE	IF	CITATIONS
1	Clearance of HIV-1 or SIV reservoirs by promotion of apoptosis and inhibition of autophagy: Targeting intracellular molecules in cure-directed strategies. <i>Journal of Leukocyte Biology</i> , 2022, 112, 1245-1259.	3.3	7
2	Identification of celastrol as a novel HIV-1 latency reversal agent by an image-based screen. <i>PLoS ONE</i> , 2021, 16, e0244771.	2.5	1
3	SARS-CoV-2 likely targets cellular PDZ proteins: a common tactic of pathogenic viruses. <i>Future Virology</i> , 2021, 16, 375-377.	1.8	9
4	Serum IgG anti-SARS-CoV-2 Binding Antibody Level Is Strongly Associated With IgA and Functional Antibody Levels in Adults Infected With SARS-CoV-2. <i>Frontiers in Immunology</i> , 2021, 12, 693462.	4.8	6
5	PACS1 is an HIV-1 cofactor that functions in Rev-mediated nuclear export of viral RNA. <i>Virology</i> , 2020, 540, 88-96.	2.4	12
6	Influenza A Virus Protein NS1 Exhibits Strain-Independent Conformational Plasticity. <i>Journal of Virology</i> , 2019, 93, .	3.4	11
7	Unexpected Mutations in HIV-1 That Confer Resistance to the Tat Inhibitor Didehydro-Cortistatin A. <i>MBio</i> , 2019, 10, .	4.1	7
8	Investigation of temporal and spatial heterogeneities of the immune responses to <i>Bordetella pertussis</i> infection in the lung and spleen of mice via analysis and modeling of dynamic microarray gene expression data. <i>Infectious Disease Modelling</i> , 2019, 4, 215-226.	1.9	1
9	Regulation of cyclin T1 during HIV replication and latency establishment in human memory CD4 T cells. <i>Virology Journal</i> , 2019, 16, 22.	3.4	8
10	Machine learning algorithms for systematic review: reducing workload in a preclinical review of animal studies and reducing human screening error. <i>Systematic Reviews</i> , 2019, 8, 23.	5.3	90
11	Roles of CDKs in RNA polymerase II transcription of the HIV-1 genome. <i>Transcription</i> , 2019, 10, 111-117.	3.1	23
12	Proteomic Profiling of a Primary CD4 <sup>+</sup> T Cell Model of HIV-1 Latency Identifies Proteins Whose Differential Expression Correlates with Reactivation of Latent HIV-1. <i>AIDS Research and Human Retroviruses</i> , 2018, 34, 103-110.	1.1	4
13	Transcriptional Control and Latency of Retroviruses. , 2018, , 199-227.		0
14	Inducible Lung Epithelial Resistance Requires Multisource Reactive Oxygen Species Generation To Protect against Viral Infections. <i>MBio</i> , 2018, 9, .	4.1	32
15	HIV-1 replication in CD4 <sup>+</sup> T cells exploits the down-regulation of antiviral NEAT1 long non-coding RNAs following T cell activation. <i>Virology</i> , 2018, 522, 193-198.	2.4	33
16	Assays for precise quantification of total (including short) and elongated HIV-1 transcripts. <i>Journal of Virological Methods</i> , 2017, 242, 1-8.	2.1	31
17	Crosstalk between histone modifications indicates that inhibition of arginine methyltransferase CARM1 activity reverses HIV latency. <i>Nucleic Acids Research</i> , 2017, 45, 9348-9360.	14.5	39
18	A scalable algorithm for structure identification of complex gene regulatory network from temporal expression data. <i>BMC Bioinformatics</i> , 2017, 18, 74.	2.6	5

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19	The HIV-1 Tat Protein: Mechanism of Action and Target for HIV-1 Cure Strategies. <i>Current Pharmaceutical Design</i> , 2017, 23, 4098-4102.	1.9	68
20	Cyclin-dependent kinases as therapeutic targets for HIV-1 infection. <i>Expert Opinion on Therapeutic Targets</i> , 2016, 20, 1453-1461.	3.4	17
21	Challenges and strategies for the eradication of the HIV reservoir. <i>Current Opinion in Immunology</i> , 2016, 42, 65-70.	5.5	54
22	Short Communication: The Broad-Spectrum Histone Deacetylase Inhibitors Vorinostat and Panobinostat Activate Latent HIV in CD4 <sup>+</sup> T Cells In Part Through Phosphorylation of the T-Loop of the CDK9 Subunit of P-TEFb. <i>AIDS Research and Human Retroviruses</i> , 2016, 32, 169-173.	1.1	21
23	Roles of microRNAs and long noncoding RNAs in human immunodeficiency virus replication. <i>Wiley Interdisciplinary Reviews RNA</i> , 2015, 6, 661-670.	6.4	21
24	Subversion of Cell Cycle Regulatory Mechanisms by HIV. <i>Cell Host and Microbe</i> , 2015, 17, 736-740.	11.0	15
25	Mining the Human Complexome Database Identifies RBM14 as an XPO1-Associated Protein Involved in HIV-1 Rev Function. <i>Journal of Virology</i> , 2015, 89, 3557-3567.	3.4	27
26	Short Communication: SAHA (Vorinostat) Induces CDK9 Thr-186 (T-Loop) Phosphorylation in Resting CD4 <sup>+</sup> T Cells: Implications for Reactivation of Latent HIV. <i>AIDS Research and Human Retroviruses</i> , 2015, 31, 137-141.	1.1	13
27	The Influenza A Virus Protein NS1 Displays Structural Polymorphism. <i>Journal of Virology</i> , 2014, 88, 4113-4122.	3.4	69
28	miR-132 enhances HIV-1 replication. <i>Virology</i> , 2013, 438, 1-4.	2.4	71
29	Reactivation of latent HIV: do all roads go through P-TEFb?. <i>Future Virology</i> , 2013, 8, 649-659.	1.8	22
30	Cyclin T1 and CDK9 T-Loop Phosphorylation Are Downregulated during Establishment of HIV-1 Latency in Primary Resting Memory CD4 <sup>+</sup> T Cells. <i>Journal of Virology</i> , 2013, 87, 1211-1220.	3.4	104
31	P-TEFb as a target to reactivate latent HIV. <i>Cell Cycle</i> , 2013, 12, 392-393.	2.6	9
32	MicroRNA-Mediated Restriction of HIV-1 in Resting CD4 <sup>+</sup> T Cells and Monocytes. <i>Viruses</i> , 2012, 4, 1390-1409.	3.3	39
33	Regulation of Cyclin T1 and HIV-1 Replication by MicroRNAs in Resting CD4 <sup>+</sup> T Lymphocytes. <i>Journal of Virology</i> , 2012, 86, 3244-3252.	3.4	153
34	Phosphatase PPM1A negatively regulates P-TEFb function in resting CD4 <sup>+</sup> T cells and inhibits HIV-1 gene expression. <i>Retrovirology</i> , 2012, 9, 52.	2.0	27
35	Identification of novel CDK9 and Cyclin T1-associated protein complexes (CCAPs) whose siRNA depletion enhances HIV-1 Tat function. <i>Retrovirology</i> , 2012, 9, 90.	2.0	19
36	Regulation of Interferon- $\beta$ by MAGI-1 and Its Interaction with Influenza A Virus NS1 Protein with ESEV PBM. <i>PLoS ONE</i> , 2012, 7, e41251.	2.5	21

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37	Making a Short Story Long: Regulation of P-TEFb and HIV-1 Transcriptional Elongation in CD4+ T Lymphocytes and Macrophages. <i>Biology</i> , 2012, 1, 94-115.	2.8	15
38	Cdk9 T-loop phosphorylation is regulated by the calcium signaling pathway. <i>Journal of Cellular Physiology</i> , 2012, 227, 609-617.	4.1	34
39	Epstein-Barr Virus BART9 miRNA Modulates LMP1 Levels and Affects Growth Rate of Nasal NK T Cell Lymphomas. <i>PLoS ONE</i> , 2011, 6, e27271.	2.5	61
40	Limited redundancy in genes regulated by Cyclin T2 and Cyclin T1. <i>BMC Research Notes</i> , 2011, 4, 260.	1.4	16
41	Emerging Theme: Cellular PDZ Proteins as Common Targets of Pathogenic Viruses. <i>Journal of Virology</i> , 2011, 85, 11544-11556.	3.4	175
42	The Avian Influenza Virus NS1 ESEV PDZ Binding Motif Associates with Dlg1 and Scribble To Disrupt Cellular Tight Junctions. <i>Journal of Virology</i> , 2011, 85, 10639-10648.	3.4	102
43	Mini ways to stop a virus: microRNAs and HIV-1 replication. <i>Future Virology</i> , 2011, 6, 209-221.	1.8	23
44	T-loop phosphorylated Cdk9 localizes to nuclear speckle domains which may serve as sites of active P-TEFb function and exchange between the Brd4 and 7SK/HEXIM1 regulatory complexes. <i>Journal of Cellular Physiology</i> , 2010, 224, 84-93.	4.1	39
45	The ESEV PDZ-Binding Motif of the Avian Influenza A Virus NS1 Protein Protects Infected Cells from Apoptosis by Directly Targeting Scribble. <i>Journal of Virology</i> , 2010, 84, 11164-11174.	3.4	90
46	55K isoform of CDK9 associates with Ku70 and is involved in DNA repair. <i>Biochemical and Biophysical Research Communications</i> , 2010, 397, 245-250.	2.1	38
47	The HIV-1 Tat Team Gets Bigger. <i>Cell Host and Microbe</i> , 2010, 7, 179-181.	11.0	2
48	Characterization of Cdk9 T-loop phosphorylation in resting and activated CD4+ T lymphocytes. <i>Journal of Leukocyte Biology</i> , 2009, 86, 1345-1350.	3.3	56
49	Dysregulation of Positive Transcription Elongation Factor b and Myocardial Hypertrophy. <i>Circulation Research</i> , 2009, 104, 1327-1329.	4.5	10
50	miR-198 Inhibits HIV-1 Gene Expression and Replication in Monocytes and Its Mechanism of Action Appears To Involve Repression of Cyclin T1. <i>PLoS Pathogens</i> , 2009, 5, e1000263.	4.7	203
51	Latently-infected CD4+ T cells are enriched for HIV-1 Tat variants with impaired transactivation activity. <i>Virology</i> , 2009, 387, 98-108.	2.4	62
52	Phosphatase PPM1A Regulates Phosphorylation of Thr-186 in the Cdk9 T-loop. <i>Journal of Biological Chemistry</i> , 2008, 283, 33578-33584.	3.4	66
53	Cyclin T1-Dependent Genes in Activated CD4+ T and Macrophage Cell Lines Appear Enriched in HIV-1 Co-Factors. <i>PLoS ONE</i> , 2008, 3, e3146.	2.5	32
54	Targeting protein-protein interactions for HIV therapeutics. <i>Future HIV Therapy</i> , 2007, 1, 369-385.	0.4	9

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55	Induction of the HIV-1 Tat co-factor cyclin T1 during monocyte differentiation is required for the regulated expression of a large portion of cellular mRNAs. <i>Retrovirology</i> , 2006, 3, 32.	2.0	25
56	Effects of prostratin on Cyclin T1/P-TEFb function and the gene expression profile in primary resting CD4+T cells. <i>Retrovirology</i> , 2006, 3, 66.	2.0	74
57	Cellular cofactors and HIV-1 infection in vivo. <i>Future Virology</i> , 2006, 1, 337-347.	1.8	2
58	Interleukin-10 inhibits HIV-1 LTR-directed gene expression in human macrophages through the induction of cyclin T1 proteolysis. <i>Virology</i> , 2006, 352, 485-492.	2.4	29
59	Cyclin T1 but not cyclin T2a is induced by a post-transcriptional mechanism in PAMP-activated monocyte-derived macrophages. <i>Journal of Leukocyte Biology</i> , 2006, 79, 388-396.	3.3	31
60	Integration of Human Immunodeficiency Virus Type 1 in Untreated Infection Occurs Preferentially within Genes. <i>Journal of Virology</i> , 2006, 80, 7765-7768.	3.4	36
61	siRNA depletion of 7SK snRNA induces apoptosis but does not affect expression of the HIV-1 LTR or P-TEFb-dependent cellular genes. <i>Journal of Cellular Physiology</i> , 2005, 205, 463-470.	4.1	33
62	Identification of LKLF-regulated genes in quiescent CD4 T lymphocytes. <i>Molecular Immunology</i> , 2005, 42, 627-641.	2.2	51
63	Human Immunodeficiency Virus Type 1 Infection Induces Cyclin T1 Expression in Macrophages. <i>Journal of Virology</i> , 2004, 78, 8114-8119.	3.4	40
64	HIV-1 infection and regulation of Tat function in macrophages. <i>International Journal of Biochemistry and Cell Biology</i> , 2004, 36, 1767-1775.	2.8	22
65	Isolation and characterization of the human DC-SIGN and DC-SIGNR promoters. <i>Gene</i> , 2003, 313, 149-159.	2.2	34
66	Increased association of 7SK snRNA with Tat cofactor P-TEFb following activation of peripheral blood lymphocytes. <i>Aids</i> , 2003, 17, 2429-2436.	2.2	32
67	Regulation of TAK / P-TEFb in CD4+ T Lymphocytes and Macrophages. <i>Current HIV Research</i> , 2003, 1, 395-404.	0.5	59
68	Transient Induction of Cyclin T1 during Human Macrophage Differentiation Regulates Human Immunodeficiency Virus Type 1 Tat Transactivation Function. <i>Journal of Virology</i> , 2002, 76, 10579-10587.	3.4	72
69	Antiapoptotic Function of Cdk9 (TAK/P-TEFb) in U937 Promonocytic Cells. <i>Journal of Virology</i> , 2001, 75, 1220-1228.	3.4	48
70	Induction of TAK (Cyclin T1/P-TEFb) in Purified Resting CD4 + T Lymphocytes by Combination of Cytokines. <i>Journal of Virology</i> , 2001, 75, 11336-11343.	3.4	99
71	Isolation and characterization of the human cyclin T1 promoter. <i>Gene</i> , 2000, 252, 39-49.	2.2	13
72	Genomic organization and characterization of promoter function of the human CDK9 gene. <i>Gene</i> , 2000, 252, 51-59.	2.2	24

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73	Tat-Associated Kinase, TAK, Activity Is Regulated by Distinct Mechanisms in Peripheral Blood Lymphocytes and Promonocytic Cell Lines. <i>Journal of Virology</i> , 1998, 72, 9881-9888.	3.4	132
74	PITALRE, the Catalytic Subunit of TAK, Is Required for Human Immunodeficiency Virus Tat Transactivation In Vivo. <i>Journal of Virology</i> , 1998, 72, 4448-4453.	3.4	111
75	RNAs Selected in vitro by the HIV-2 Tat Protein. <i>Journal of Biomedical Science</i> , 1997, 4, 28-34.	7.0	0
76	RNAs selected in vitro by the HIV-2 tat protein. <i>Journal of Biomedical Science</i> , 1997, 4, 28-34.	7.0	7
77	Wild-Type and Transactivation-Defective Mutants of Human Immunodeficiency Virus Type 1 Tat Protein Bind Human TATA-Binding Protein In Vitro. <i>Journal of Acquired Immune Deficiency Syndromes</i> , 1996, 12, 128-138.	0.3	5
78	HIV-1 Tat protein is able to efficiently transactivate the HIV-2 LTR through a TAR RNA element lacking both dinucleotide bulge binding sites. <i>Virology</i> , 1995, 206, 673-678.	2.4	7
79	Exon2 of HIV-2 Tat Contributes to transactivation of the HIV-2 LTR by increasing binding affinity to HIV-2 TAR RNA. <i>Nucleic Acids Research</i> , 1994, 22, 4405-4413.	14.5	26
80	Functional Significance of the Dinucleotide Bulge in Stem-Loop1 and Stem-Loop2 of HIV-2 TAR RNA. <i>Virology</i> , 1994, 202, 202-211.	2.4	21
81	Specific Interaction of the Human Immunodeficiency Virus Tat Proteins with a Cellular Protein Kinase. <i>Virology</i> , 1993, 197, 601-608.	2.4	143
82	Tat protein of human immunodeficiency virus type 1 is a monomer when expressed in mammalian cells. <i>Virology</i> , 1991, 185, 451-454.	2.4	28
83	Regulation of HIV-1 Gene Expression by the Tat Protein and the TAR Region. , 1991, , 93-105.		0
84	HIV-1 Tat protein increases transcriptional initiation and stabilizes elongation. <i>Cell</i> , 1989, 59, 283-292.	28.9	653
85	Transcriptional but not translational regulation of HIV-1 by the tat gene product. <i>Nature</i> , 1988, 332, 551-553.	27.8	192