

# Shunbin Ning

## List of Publications by Year in descending order

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Version: 2024-02-01

72  
papers

2,347  
citations

293460

24  
h-index

263392

45  
g-index

73  
all docs

73  
docs citations

73  
times ranked

3823  
citing authors

#	ARTICLE	IF	CITATIONS
1	TRF2 inhibition rather than telomerase disruption drives CD4T cell dysfunction during chronic viral infection. <i>Journal of Cell Science</i> , 2022, 135, .	1.2	4
2	TRIMming Type I Interferon-Mediated Innate Immune Response in Antiviral and Antitumor Defense. <i>Viruses</i> , 2021, 13, 279.	1.5	18
3	Long Non-coding RNA GAS5 Regulates T Cell Functions via miR21-Mediated Signaling in People Living With HIV. <i>Frontiers in Immunology</i> , 2021, 12, 601298.	2.2	24
4	Algorithm-Based Meta-Analysis Reveals the Mechanistic Interaction of the Tumor Suppressor LIMD1 With Non-Small-Cell Lung Carcinoma. <i>Frontiers in Oncology</i> , 2021, 11, 632638.	1.3	3
5	Blockade of SARS-CoV-2 spike protein-mediated cell-cell fusion using COVID-19 convalescent plasma. <i>Scientific Reports</i> , 2021, 11, 5558.	1.6	19
6	Long Noncoding RNA RUNXOR Promotes Myeloid-Derived Suppressor Cell Expansion and Functions via Enhancing Immunosuppressive Molecule Expressions during Latent HIV Infection. <i>Journal of Immunology</i> , 2021, 206, 2052-2060.	0.4	7
7	Mitochondrial Functions Are Compromised in CD4 T Cells From ART-Controlled PLHIV. <i>Frontiers in Immunology</i> , 2021, 12, 658420.	2.2	20
8	Immune Activation Induces Telomeric DNA Damage and Promotes Short-Lived Effector T Cell Differentiation in Chronic HCV Infection. <i>Hepatology</i> , 2021, 74, 2380-2394.	3.6	11
9	The Ubiquitin Sensor and Adaptor Protein p62 Mediates Signal Transduction of a Viral Oncogenic Pathway. <i>MBio</i> , 2021, 12, e0109721.	1.8	8
10	SARS-CoV-2 specific memory T cell epitopes identified in COVID-19-recovered subjects. <i>Virus Research</i> , 2021, 304, 198508.	1.1	31
11	New Look of EBV LMP1 Signaling Landscape. <i>Cancers</i> , 2021, 13, 5451.	1.7	23
12	How Oncogenic Viruses Exploit p62-Mediated Selective Autophagy for Cancer Development. <i>Annals of Immunology &amp; Immunotherapy</i> , 2021, 3, .	0.1	0
13	Selective oxidative stress induces dual damage to telomeres and mitochondria in human T cells. <i>Aging Cell</i> , 2021, 20, e13513.	3.0	39
14	Oxidative Stress Induces Mitochondrial Compromise in CD4 T Cells From Chronically HCV-Infected Individuals. <i>Frontiers in Immunology</i> , 2021, 12, 760707.	2.2	5
15	Telomeric injury by KML001 in human T cells induces mitochondrial dysfunction through the p53-PGC-1 pathway. <i>Cell Death and Disease</i> , 2020, 11, 1030.	2.7	23
16	Telomere and ATM Dynamics in CD4 T-Cell Depletion in Active and Virus-Suppressed HIV Infections. <i>Journal of Virology</i> , 2020, 94, .	1.5	9
17	HCV-Associated Exosomes Upregulate RUNXOR and RUNX1 Expressions to Promote MDSC Expansion and Suppressive Functions through STAT3-miR124 Axis. <i>Cells</i> , 2020, 9, 2715.	1.8	33
18	Inhibition of topoisomerase IIA (Top2 $\alpha$ ) induces telomeric DNA damage and T cell dysfunction during chronic viral infection. <i>Cell Death and Disease</i> , 2020, 11, 196.	2.7	21

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19	A Matter of Life or Death: Productively Infected and Bystander CD4 T Cells in Early HIV Infection. <i>Frontiers in Immunology</i> , 2020, 11, 626431.	2.2	18
20	LncRNA HOTAIRM1 promotes MDSC expansion and suppressive functions through the HOXA1-miR124 axis during HCV infection. <i>Scientific Reports</i> , 2020, 10, 22033.	1.6	19
21	Long noncoding RNA HOTAIRM1 promotes myeloid-derived suppressor cell expansion and suppressive functions through up-regulating HOXA1 expression during latent HIV infection. <i>Aids</i> , 2020, 34, 2211-2221.	1.0	16
22	Topological DNA damage, telomere attrition and T cell senescence during chronic viral infections. <i>Immunity and Ageing</i> , 2019, 16, 12.	1.8	26
23	Disruption of Telomere Integrity and DNA Repair Machineries by KML001 Induces T Cell Senescence, Apoptosis, and Cellular Dysfunctions. <i>Frontiers in Immunology</i> , 2019, 10, 1152.	2.2	26
24	p62-mediated Selective autophagy endows virus-transformed cells with insusceptibility to DNA damage under oxidative stress. <i>PLoS Pathogens</i> , 2019, 15, e1007541.	2.1	42
25	ATM Deficiency Accelerates DNA Damage, Telomere Erosion, and Premature T Cell Aging in HIV-Infected Individuals on Antiretroviral Therapy. <i>Frontiers in Immunology</i> , 2019, 10, 2531.	2.2	27
26	The Multifunctional Protein p62 and Its Mechanistic Roles in Cancers. <i>Current Cancer Drug Targets</i> , 2019, 19, 468-478.	0.8	22
27	Insufficiency of DNA repair enzyme ATM promotes naive CD4 T-cell loss in chronic hepatitis C virus infection. <i>Cell Discovery</i> , 2018, 4, 16.	3.1	40
28	HCV-associated exosomes promote myeloid-derived suppressor cell expansion via inhibiting miR-124 to regulate T follicular cell differentiation and function. <i>Cell Discovery</i> , 2018, 4, 51.	3.1	34
29	Inhibition of TRF2 accelerates telomere attrition and DNA damage in naïve CD4 T cells during HCV infection. <i>Cell Death and Disease</i> , 2018, 9, 900.	2.7	27
30	LIMD1 is induced by and required for LMP1 signaling, and protects EBV-transformed cells from DNA damage-induced cell death. <i>Oncotarget</i> , 2018, 9, 6282-6297.	0.8	17
31	The Linear Ubiquitin Assembly Complex Modulates Latent Membrane Protein 1 Activation of NF- $\kappa$ B and Interferon Regulatory Factor 7. <i>Journal of Virology</i> , 2017, 91, .	1.5	23
32	Decline of miR-124 in myeloid cells promotes regulatory T cell development in hepatitis C virus infection. <i>Immunology</i> , 2017, 150, 213-220.	2.0	19
33	LMP1 signaling pathway activates IRF4 in latent EBV infection and a positive circuit between PI3K and Src is required. <i>Oncogene</i> , 2017, 36, 2265-2274.	2.6	25
34	Identification of <i>KANSARL</i> as the first cancer predisposition fusion gene specific to the population of European ancestry origin. <i>Oncotarget</i> , 2017, 8, 50594-50607.	0.8	24
35	“Toll-free” pathways for production of type I interferons. <i>AIMS Allergy and Immunology</i> , 2017, 1, 143-163.	0.3	9
36	Identification of PP1 as the First Phosphatase for IRF7. <i>Journal of Cell Signaling</i> , 2017, 02, .	0.3	1

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37	Hepatitis C virus-induced myeloid-derived suppressor cells regulate T cell differentiation and function via the signal transducer and activator of transcription 3 pathway. <i>Immunology</i> , 2016, 148, 377-386.	2.0	47
38	HCV-induced miR146a controls SOCS1/STAT3 and cytokine expression in monocytes to promote regulatory T cell development. <i>Journal of Viral Hepatitis</i> , 2016, 23, 755-766.	1.0	20
39	Protein phosphatase 1 abrogates IRF7-mediated type I IFN response in antiviral immunity. <i>European Journal of Immunology</i> , 2016, 46, 2409-2419.	1.6	34
40	Protection of CD4+ T cells from hepatitis C virus infection-associated senescence via miR-181a/Sirt1 pathway. <i>Journal of Leukocyte Biology</i> , 2016, 100, 1201-1211.	1.5	25
41	Expansion of myeloid-derived suppressor cells promotes differentiation of regulatory T cells in HIV-1+ individuals. <i>Aids</i> , 2016, 30, 1521-1531.	1.0	64
42	MicroRNA regulation of viral immunity, latency, and carcinogenesis of selected tumor viruses and HIV. <i>Reviews in Medical Virology</i> , 2015, 25, 320-341.	3.9	21
43	Human DNA Exonuclease TREX1 Is Also an Exoribonuclease That Acts on Single-stranded RNA. <i>Journal of Biological Chemistry</i> , 2015, 290, 13344-13353.	1.6	31
44	Gene Expression Profiling Identifies IRF4-Associated Molecular Signatures in Hematological Malignancies. <i>PLoS ONE</i> , 2014, 9, e106788.	1.1	34
45	Interferon Regulatory Factor 4 Is Activated through c-Src-Mediated Tyrosine Phosphorylation in Virus-Transformed Cells. <i>Journal of Virology</i> , 2013, 87, 9672-9679.	1.5	16
46	IRF4 as an Oncogenic Biomarker for Hematological Malignancies. <i>Journal of Oncobiomarkers</i> , 2013, 1, .	0.3	6
47	IRF7: activation, regulation, modification and function. <i>Genes and Immunity</i> , 2011, 12, 399-414.	2.2	428
48	Innate immune modulation in EBV infection. <i>Herpesviridae</i> , 2011, 2, 1.	2.7	60
49	Oncogenic IRFs Provide a Survival Advantage for Epstein-Barr Virus- or Human T-Cell Leukemia Virus Type 1-Transformed Cells through Induction of BIC Expression. <i>Journal of Virology</i> , 2011, 85, 8328-8337.	1.5	50
50	The A20 Deubiquitinase Activity Negatively Regulates LMP1 Activation of IRF7. <i>Journal of Virology</i> , 2010, 84, 6130-6138.	1.5	63
51	PS2-61 Transcriptional regulation of miR-155 by IRFs in antiviral immunity and viral tumors. <i>Cytokine</i> , 2010, 52, 63.	1.4	0
52	The Epstein-Barr Virus (EBV) Deubiquitinating Enzyme BPLF1 Reduces EBV Ribonucleotide Reductase Activity. <i>Journal of Virology</i> , 2009, 83, 4345-4353.	1.5	63
53	TRAF6 and the Three C-Terminal Lysine Sites on IRF7 Are Required for Its Ubiquitination-Mediated Activation by the Tumor Necrosis Factor Receptor Family Member Latent Membrane Protein 1. <i>Molecular and Cellular Biology</i> , 2008, 28, 6536-6546.	1.1	94
54	Interferon Regulatory Factor 7 Is Activated by a Viral Oncoprotein through RIP-Dependent Ubiquitination. <i>Molecular and Cellular Biology</i> , 2007, 27, 2910-2918.	1.1	69

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55	Regulation of the Transcriptional Activity of the IRF7 Promoter by a Pathway Independent of Interferon Signaling. <i>Journal of Biological Chemistry</i> , 2005, 280, 12262-12270.	1.6	77
56	Interferon Regulatory Factor 7 Is Negatively Regulated by the Epstein-Barr Virus Immediate-Early Gene, BZLF-1. <i>Journal of Virology</i> , 2005, 79, 10040-10052.	1.5	98
57	Interferon Regulatory Factor 5 Represses Expression of the Epstein-Barr Virus Oncoprotein LMP1: Braking of the IRF7/LMP1 Regulatory Circuit. <i>Journal of Virology</i> , 2005, 79, 11671-11676.	1.5	41
58	Interferon Regulatory Factor 7 Regulates Expression of Epstein-Barr Virus Latent Membrane Protein 1: a Regulatory Circuit. <i>Journal of Virology</i> , 2003, 77, 9359-9368.	1.5	88
59	Identification of programmed cell death in situ in individual plant cells in vivo using a chromosome preparation technique. <i>Journal of Experimental Botany</i> , 2002, 53, 651-658.	2.4	31
60	Characterization of the early stages of programmed cell death in maize root cells by using comet assay and the combination of cell electrophoresis with annexin binding. <i>Electrophoresis</i> , 2002, 23, 2096.	1.3	27
61	Salt stress induces programmed cell death in prokaryotic organism <i>Anabaena</i> . <i>Journal of Applied Microbiology</i> , 2002, 93, 15-28.	1.4	111
62	Determination of copy number for 5S rDNA and centromeric sequence RCS2 in rice by Fiber-FISH. <i>Science Bulletin</i> , 2002, 47, 214.	1.7	4
63	Apoptotic Cell Death and Cellular Surface Negative Charge Increase in Maize Roots Exposed to Cytotoxic Stresses. <i>Annals of Botany</i> , 2001, 87, 575-583.	1.4	15
64	FISH analysis of the integration patterns in transgenic rice co-transformed by microprojectile bombardment. <i>Science Bulletin</i> , 2001, 46, 1965-1968.	1.7	2
65	Comparative genome research between maize and rice using genomic in situ hybridization. <i>Science Bulletin</i> , 2001, 46, 656-658.	1.7	2
66	Physical location of rice <i>Gm-6</i> , <i>Pi-5(t)</i> genes in <i>O. officinalis</i> with BAC-FISH. <i>Science Bulletin</i> , 2001, 46, 659-661.	1.7	8
67	Detection of Alien Genes and Analysis of their Integration Position in Transgenic Rice by Fluorescence <i>in situ</i> Hybridization. <i>Breeding Science</i> , 2001, 51, 279-283.	0.9	0
68	Mammalian Apoptosis-associated Genes <i>c-myc</i> and <i>p53</i> in Maize. Homologs and Their Locations. <i>Cytologia</i> , 2000, 65, 261-270.	0.2	0
69	Maize <i>nacl</i> and <i>cld</i> genes map to chromosome arms 10L and 2S, and to 4L and 5L, respectively. <i>Chromosome Research</i> , 2000, 8, 273-273.	1.0	0
70	An NMR study of the structural basis of the wide range of pharmacological functions of acetylsalicylic acid. <i>IUBMB Life</i> , 1999, 47, 665-671.	1.5	2
71	A novel method for in situ detection of apoptotic cell death in plants. <i>Science Bulletin</i> , 1999, 44, 1014-1017.	1.7	2
72	Bioinformatics-Driven Identification of <i>p62</i> as A Crucial Oncogene in Liver Cancer. <i>Frontiers in Oncology</i> , 0, 12, .	1.3	1