Hualin Wang

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Taxonomy of the order Bunyavirales: update 2019. Archives of Virology, 2019, 164, 1949-1965.	0.9	285
2	The sequence of the Helicoverpa armigera single nucleocapsid nucleopolyhedrovirus genome. Journal of General Virology, 2001, 82, 241-257.	1.3	200
3	Proteomics of the <i>Autographa californica</i> Nucleopolyhedrovirus Budded Virions. Journal of Virology, 2010, 84, 7233-7242.	1.5	150
4	Disruption of Type I Interferon Signaling by the Nonstructural Protein of Severe Fever with Thrombocytopenia Syndrome Virus via the Hijacking of STAT2 and STAT1 into Inclusion Bodies. Journal of Virology, 2015, 89, 4227-4236.	1.5	106
5	Viral suppression of innate immunity via spatial isolation of TBK1/IKKε from mitochondrial antiviral platform. Journal of Molecular Cell Biology, 2014, 6, 324-337.	1.5	92
6	Comparative Proteomics Reveal Fundamental Structural and Functional Differences between the Two Progeny Phenotypes of a Baculovirus. Journal of Virology, 2013, 87, 829-839.	1.5	87
7	A novel tick-borne phlebovirus, closely related to severe fever with thrombocytopenia syndrome virus and Heartland virus, is a potential pathogen. Emerging Microbes and Infections, 2018, 7, 1-14.	3.0	78
8	Proteomics Analysis of Helicoverpa armigera Single Nucleocapsid Nucleopolyhedrovirus Identified Two New Occlusion-Derived Virus-Associated Proteins, HA44 and HA100. Journal of Virology, 2007, 81, 9377-9385.	1.5	77
9	SARS-CoV-2 nsp1: Bioinformatics, Potential Structural and Functional Features, and Implications for Drug/Vaccine Designs. Frontiers in Microbiology, 2020, 11, 587317.	1.5	60
10	Functional studies of per os infectivity factors of Helicoverpa armigera single nucleocapsid nucleopolyhedrovirus. Journal of General Virology, 2008, 89, 2331-2338.	1.3	55
11	Migration, recombination, and reassortment are involved in the evolution of severe fever with thrombocytopenia syndrome bunyavirus. Infection, Genetics and Evolution, 2017, 47, 109-117.	1.0	54
12	lmmune evasion of SARS-CoV-2 from interferon antiviral system. Computational and Structural Biotechnology Journal, 2021, 19, 4217-4225.	1.9	49
13	<i>Autographa californica</i> Multicapsid Nucleopolyhedrovirus Efficiently Infects Sf9 Cells and Transduces Mammalian Cells via Direct Fusion with the Plasma Membrane at Low pH. Journal of Virology, 2010, 84, 5351-5359.	1.5	48
14	Heartland virus NSs protein disrupts host defenses by blocking the TBK1 kinase–IRF3 transcription factor interaction and signaling required for interferon induction. Journal of Biological Chemistry, 2017, 292, 16722-16733.	1.6	46
15	Phylogenetic analysis revealed the central roles of two African countries in the evolution and worldwide spread of Zika virus. Virologica Sinica, 2016, 31, 118-130.	1.2	45
16	Function, oligomerization and N-linked glycosylation of the Helicoverpa armigera single nucleopolyhedrovirus envelope fusion protein. Journal of General Virology, 2006, 87, 839-846.	1.3	44
17	The nucleoprotein of severe fever with thrombocytopenia syndrome virus processes a stable hexameric ring to facilitate RNA encapsidation. Protein and Cell, 2013, 4, 445-455.	4.8	44
18	Zika Virus Baculovirus-Expressed Virus-Like Particles Induce Neutralizing Antibodies in Mice. Virologica Sinica, 2018, 33, 213-226.	1.2	43

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19	Construction and Rescue of a Functional Synthetic Baculovirus. ACS Synthetic Biology, 2017, 6, 1393-1402.	1.9	40
20	A RIG-l–like receptor directs antiviral responses to a bunyavirus and is antagonized by virus-induced blockade of TRIM25-mediated ubiquitination. Journal of Biological Chemistry, 2020, 295, 9691-9711.	1.6	39
21	The F-Like Protein Ac23 Enhances the Infectivity of the Budded Virus of <i>gp64</i> -Null <i>Autographa californica</i> Multinucleocapsid Nucleopolyhedrovirus Pseudotyped with Baculovirus Envelope Fusion Protein F. Journal of Virology, 2008, 82, 9800-9804.	1.5	38
22	Crystal Structure of the Core Region of Hantavirus Nucleocapsid Protein Reveals the Mechanism for Ribonucleoprotein Complex Formation. Journal of Virology, 2016, 90, 1048-1061.	1.5	35
23	Pathologic Studies of Fatal Encephalomyelitis in Children Caused by Enterovirus 71. American Journal of Clinical Pathology, 2016, 146, 95-106.	0.4	34
24	Open reading frame 132 of Heliocoverpa armigera nucleopolyhedrovirus encodes a functional per os infectivity factor (PIF-2). Journal of General Virology, 2006, 87, 2563-2569.	1.3	32
25	Open reading frame 94 of Helicoverpa armigera single nucleocapsid nucleopolyhedrovirus encodes a novel conserved occlusion-derived virion protein, ODV-EC43. Journal of General Virology, 2003, 84, 3021-3027.	1.3	31
26	Virus like particle-based vaccines against emerging infectious disease viruses. Virologica Sinica, 2016, 31, 279-287.	1.2	31
27	Singleâ€Particle Tracking Reveals the Sequential Entry Process of the Bunyavirus Severe Fever with Thrombocytopenia Syndrome Virus. Small, 2019, 15, e1803788.	5.2	31
28	Heartland virus antagonizes type I and III interferon antiviral signaling by inhibiting phosphorylation and nuclear translocation of STAT2 and STAT1. Journal of Biological Chemistry, 2019, 294, 9503-9517.	1.6	30
29	Baculovirus <i>Per Os</i> Infectivity Factor Complex: Components and Assembly. Journal of Virology, 2019, 93, .	1.5	29
30	Reassortment and migration analysis of Crimean–Congo haemorrhagic fever virus. Journal of General Virology, 2013, 94, 2536-2548.	1.3	27
31	Unraveling the Entry Mechanism of Baculoviruses and Its Evolutionary Implications. Journal of Virology, 2014, 88, 2301-2311.	1.5	27
32	Interferon-Î ³ -Directed Inhibition of a Novel High-Pathogenic Phlebovirus and Viral Antagonism of the Antiviral Signaling by Targeting STAT1. Frontiers in Immunology, 2019, 10, 1182.	2.2	26
33	Functional analysis of FP25K of Helicoverpa armigera single nucleocapsid nucleopolyhedrovirus. Journal of General Virology, 2005, 86, 2439-2444.	1.3	25
34	Genomic and transcriptional analyses of novel parvoviruses identified from dead peafowl. Virology, 2020, 539, 80-91.	1.1	25
35	Quantitative Proteomic Analysis Reveals Unfolded-Protein Response Involved in Severe Fever with Thrombocytopenia Syndrome Virus Infection. Journal of Virology, 2019, 93, .	1.5	24
36	Genome Sequence and Analysis of Buzura suppressaria Nucleopolyhedrovirus: A Group II Alphabaculovirus. PLoS ONE, 2014, 9, e86450.	1,1	21

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37	Per os infectivity factors: a complicated and evolutionarily conserved entry machinery of baculovirus. Science China Life Sciences, 2017, 60, 806-815.	2.3	21
38	First case of laboratory-confirmed severe fever with thrombocytopenia syndrome disease revealed the risk of SFTSV infection in Xinjiang, China. Emerging Microbes and Infections, 2019, 8, 1122-1125.	3.0	21
39	Host restriction of emerging high-pathogenic bunyaviruses via MOV10 by targeting viral nucleoprotein and blocking ribonucleoprotein assembly. PLoS Pathogens, 2020, 16, e1009129.	2.1	21
40	Characterization and phylogenetic analysis of the chitinase gene from the Helicoverpa armigera single nucleocapsid nucleopolyhedrovirus. Virus Research, 2004, 100, 179-189.	1.1	20
41	Open reading frame Bm21 of Bombyx mori nucleopolyhedrovirus is not essential for virus replication in vitro, but its deletion extends the median survival time of infected larvae. Journal of General Virology, 2008, 89, 922-930.	1.3	19
42	Crimean-Congo Hemorrhagic Fever Virus: Current Advances and Future Prospects of Antiviral Strategies. Viruses, 2021, 13, 1195.	1.5	19
43	The Host Specificities of Baculovirus per os Infectivity Factors. PLoS ONE, 2016, 11, e0159862.	1.1	19
44	The F protein of Helicoverpa armigera single nucleopolyhedrovirus can be substituted functionally with its homologue from Spodoptera exigua multiple nucleopolyhedrovirus. Journal of General Virology, 2008, 89, 791-798.	1.3	18
45	Angiotensin-converting enzyme 2 (ACE2) from raccoon dog can serve as an efficient receptor for the spike protein of severe acute respiratory syndrome coronavirus. Journal of General Virology, 2009, 90, 2695-2703.	1.3	18
46	A novel glycoprotein D-specific monoclonal antibody neutralizes herpes simplex virus. Antiviral Research, 2017, 147, 131-141.	1.9	18
47	The Functional Oligomeric State of Tegument Protein GP41 Is Essential for Baculovirus Budded Virion and Occlusion-Derived Virion Assembly. Journal of Virology, 2018, 92, .	1.5	18
48	The roles of ebolavirus glycoproteins in viral pathogenesis. Virologica Sinica, 2017, 32, 3-15.	1.2	17
49	Viromes and surveys of RNA viruses in camel-derived ticks revealing transmission patterns of novel tick-borne viral pathogens in Kenya. Emerging Microbes and Infections, 2021, 10, 1975-1987.	3.0	17
50	The Complete Genome of a New Betabaculovirus from Clostera anastomosis. PLoS ONE, 2015, 10, e0132792.	1.1	15
51	Characterization of two monoclonal antibodies, 38F10 and 44D11, against the major envelope fusion protein of Helicoverpa armigera nucleopolyhedrovirus. Virologica Sinica, 2016, 31, 490-499.	1.2	14
52	Interactome profiling reveals interaction of SARS-CoV-2 NSP13 with host factor STAT1 to suppress interferon signaling. Journal of Molecular Cell Biology, 2021, 13, 760-762.	1.5	14
53	Evidence of Human Exposure to Tamdy Virus, Northwest China. Emerging Infectious Diseases, 2021, 27, 3166-3170.	2.0	14
54	Baculovirus ODV-E66 degrades larval peritrophic membrane to facilitate baculovirus oral infection. Virology, 2019, 537, 157-164.	1.1	13

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55	The Nonstructural Protein of Guertu Virus Disrupts Host Defenses by Blocking Antiviral Interferon Induction and Action. ACS Infectious Diseases, 2020, 6, 857-870.	1.8	13
56	Nucleotide sequence and transcriptional analysis of a putative basic DNA-binding protein of Helicoverpa armigera nucleopolyhedrovirus. Virus Genes, 2001, 22, 113-120.	0.7	12
57	Helicoverpa armigera nucleopolyhedrovirus occlusion-derived virus-associated protein, HA100, affects oral infectivity in vivo but not virus replication in vitro. Journal of General Virology, 2011, 92, 1324-1331.	1.3	12
58	Three Conserved Regions in Baculovirus Sulfhydryl Oxidase P33 Are Critical for Enzymatic Activity and Function. Journal of Virology, 2017, 91, .	1.5	12
59	The Major Hurdle for Effective Baculovirus Transduction into Mammalian Cells Is Passing Early Endosomes. Journal of Virology, 2019, 93, .	1.5	12
60	Genome Sequencing and Analysis of Catopsilia pomona nucleopolyhedrovirus: A Distinct Species in Group I Alphabaculovirus. PLoS ONE, 2016, 11, e0155134.	1.1	11
61	Resistant mutations and quasispecies complexity of hepatitis B virus during telbivudine treatment. Journal of General Virology, 2015, 96, 3302-3312.	1.3	11
62	The group I alphabaculovirus-specific protein, AC5, is a novel component of the occlusion body but is not associated with ODVs or the PIF complex. Journal of General Virology, 2018, 99, 585-595.	1.3	11
63	Glycoprotein E of the Japanese encephalitis virus forms virus-like particles and induces syncytia when expressed by a baculovirus. Journal of General Virology, 2015, 96, 1006-1014.	1.3	10
64	Combinatorial Minigenome Systems for Emerging Banyangviruses Reveal Viral Reassortment Potential and Importance of a Protruding Nucleotide in Genome "Panhandle―for Promoter Activity and Reassortment. Frontiers in Microbiology, 2020, 11, 599.	1.5	10
65	Novel SFTSV Phylogeny Reveals New Reassortment Events and Migration Routes. Virologica Sinica, 2021, 36, 300-310.	1.2	10
66	Genomic Sequencing and Analysis of Sucra jujuba Nucleopolyhedrovirus. PLoS ONE, 2014, 9, e110023.	1.1	10
67	Deletion of a Helicoverpa armigera nucleopolyhedrovirus gene encoding a virion structural protein (ORF107) increases the budded virion titre and reduces in vivo infectivity. Journal of General Virology, 2007, 88, 3307-3316.	1.3	9
68	Mutational and functional analysis of N-linked glycosylation of envelope fusion protein F of Helicoverpa armigera nucleopolyhedrovirus. Journal of General Virology, 2016, 97, 988-999.	1.3	9
69	Animal Model of Severe Fever With Thrombocytopenia Syndrome Virus Infection. Frontiers in Microbiology, 2021, 12, 797189.	1.5	9
70	Betabaculovirus F proteins showed different efficiencies when rescuing the infectivity of gp64-null Autographa californica nucleopolyhedrovirus. Virology, 2013, 436, 59-66.	1.1	8
71	Characterization of the viral fibroblast growth factor homolog of Helicoverpa armigera single nucleopolyhedrovirus. Virologica Sinica, 2016, 31, 240-248.	1.2	8
72	Improving Baculovirus Transduction of Mammalian Cells by Incorporation of Thogotovirus Glycoproteins. Virologica Sinica, 2019, 34, 454-466.	1.2	8

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73	Establishment of a Reverse Genetic System of Severe Fever with Thrombocytopenia Syndrome Virus Based on a C4 Strain. Virologica Sinica, 2021, 36, 958-967.	1.2	8
74	Recent Advances in Bunyavirus Reverse Genetics Research: Systems Development, Applications, and Future Perspectives. Frontiers in Microbiology, 2021, 12, 771934.	1.5	8
75	Incorporation of GP64 into Helicoverpa armigera nucleopolyhedrovirus enhances virus infectivity in vivo and in vitro. Journal of General Virology, 2012, 93, 2705-2711.	1.3	7
76	Host factor heat-shock protein 90 contributes to baculovirus budded virus morphogenesis via facilitating nuclear actin polymerization. Virology, 2019, 535, 200-209.	1.1	7
77	Systematic Analysis of 42 Autographa Californica Multiple Nucleopolyhedrovirus Genes Identifies An Additional Six Genes Involved in the Production of Infectious Budded Virus. Virologica Sinica, 2021, 36, 762-773.	1.2	7
78	The FP25K Acts as a Negative Factor for the Infectivity of AcMNPV Budded Virus. PLoS ONE, 2015, 10, e0128471.	1.1	6
79	Immunological detection of serum antibodies in pediatric medical workers exposed to varying levels of SARS-CoV-2. Journal of Infection, 2021, 82, 159-198.	1.7	6
80	Construction and Characterization of a Novel Bacmid AcBac-Syn Based on a Synthesized Baculovirus Genome. Virologica Sinica, 2021, 36, 1566-1574.	1.2	6
81	Genome analysis of a novel Group I alphabaculovirus obtained from Oxyplax ochracea. PLoS ONE, 2018, 13, e0192279.	1.1	6
82	Functional studies of per os infectivity factor 3 of Helicoverpa armigera nucleopolyhedrovirus. Journal of General Virology, 2012, 93, 374-382.	1.3	5
83	P33 of Helicoverpa armigera single nucleocapsid nucleopolyhedrovirus is a functional homolog of AcP33. Virologica Sinica, 2016, 31, 346-349.	1.2	5
84	Ebola virus mucin-like glycoprotein (Emuc) induces remarkable acute inflammation and tissue injury: evidence for Emuc pathogenicity in vivo. Protein and Cell, 2017, 9, 389-393.	4.8	5
85	Functional Characterization of the Group I Alphabaculovirus Specific Gene ac73. Virologica Sinica, 2019, 34, 701-711.	1.2	5
86	The cysteine-rich region of a baculovirus VP91 protein contributes to the morphogenesis of occlusion bodies. Virology, 2019, 535, 144-153.	1.1	5
87	<i>Per Os</i> Infectivity Factor 5 Identified as a Substrate of P33 in the Baculoviral Disulfide Bond Formation Pathway. Journal of Virology, 2020, 94, .	1.5	5
88	An in vitro recombination-based reverse genetic system for rapid mutagenesis of structural genes of the Japanese encephalitis virus. Virologica Sinica, 2015, 30, 354-362.	1.2	4
89	HearNPV Pseudotyped with PIF1, 2, and 3 from MabrNPV: Infectivity and Complex Stability. Virologica Sinica, 2018, 33, 187-196.	1.2	4
90	Establishment of Baculovirus-Expressed VLPs Induced Syncytial Formation Assay for Flavivirus Antiviral Screening. Viruses, 2018, 10, 365.	1.5	4

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91	Host AAA+ ATPase TER94 Plays Critical Roles in Building the Baculovirus Viral Replication Factory and Virion Morphogenesis. Journal of Virology, 2020, 94, .	1.5	4
92	Systematic analysis of nuclear localization of Autographa californica multiple nucleopolyhedrovirus proteins. Journal of General Virology, 2021, 102, .	1.3	4
93	Non-structural Proteins of Severe Fever With Thrombocytopenia Syndrome Virus Suppress RNA Synthesis in a Transcriptionally Active cDNA-Derived Viral RNA Synthesis System. Frontiers in Microbiology, 2021, 12, 709517.	1.5	4
94	Genome Characteristics of the Cyclophragma Undans Nucleopolyhedrovirus: A Distinct Species in Group I of Alphabaculovirus. Virologica Sinica, 2018, 33, 359-368.	1.2	3
95	Genome Analysis of a Novel Clade II.b Alphabaculovirus Obtained from Artaxa digramma. Viruses, 2019, 11, 925.	1.5	3
96	Identification, Isolation, and Characterization of an Ectromelia Virus New Strain from an Experimental Mouse. Virologica Sinica, 2021, 36, 155-158.	1.2	3
97	Identification and functional analysis of inter-subunit disulfide bonds of the F protein of Helicoverpa armigera nucleopolyhedrovirus. Journal of General Virology, 2014, 95, 2820-2830.	1.3	2
98	Mitoâ€docking: A Novel In Vivo Method to Detect Protein–Protein Interactions. Small Methods, 2019, 3, 1900010.	4.6	2
99	Deltabaculoviruses encode a functional type I budded virus envelope fusion protein. Journal of General Virology, 2017, 98, 847-852.	1.3	2
100	Development of Multi-analyte Suspension Assay for Simultaneously Efficient Detection of Avian Influenza Virus A Subtypes. Virologica Sinica, 2018, 33, 111-115.	1.2	1
101	Bunyaviruses: Singleâ€Particle Tracking Reveals the Sequential Entry Process of the Bunyavirus Severe Fever with Thrombocytopenia Syndrome Virus (Small 6/2019). Small, 2019, 15, 1970032.	5.2	1