

Xuan Jiang

List of Publications by Year in descending order

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Version: 2024-02-01

23
papers

955
citations

858243

12
h-index

759306

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26
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times ranked

1874
citing authors

#	ARTICLE	IF	CITATIONS
1	iTRAQ-Based Proteomics Analysis of Human Cytomegalovirus Latency and Reactivation in T98G Cells. <i>Journal of Virology</i> , 2022, 96, JVI0147621.	1.5	4
2	Human Cytomegalovirus Hijacks WD Repeat Domain 11 for Virion Assembly Compartment Formation and Virion Morphogenesis. <i>Journal of Virology</i> , 2022, 96, JVI0182721.	1.5	4
3	A congenital CMV infection model for follow-up studies of neurodevelopmental disorders, neuroimaging abnormalities, and treatment. <i>JCI Insight</i> , 2022, 7, .	2.3	17
4	Localization of the WD Repeat-Containing Protein 5 to the Virion Assembly Compartment Facilitates Human Cytomegalovirus Assembly. <i>Journal of Virology</i> , 2021, 95, .	1.5	3
5	Hearing Loss Caused by HCMV Infection through Regulating the Wnt and Notch Signaling Pathways. <i>Viruses</i> , 2021, 13, 623.	1.5	7
6	Establishing an Animal Model of Cytomegalovirus Keratouveitis in Rats: Broad Infection of Anterior Segment Tissue by Cytomegalovirus. , 2021, 62, 22.		6
7	Pathogenic Effects and Pathogenesis Processes in Vitro & in Vivo in Murine Cytomegalovirus Infected Rat Corneal Endothelial Cells. <i>Ocular Immunology and Inflammation</i> , 2020, , 1-12.	1.0	3
8	The Golgi Apparatus May Be a Potential Therapeutic Target for Apoptosis-Related Neurological Diseases. <i>Frontiers in Cell and Developmental Biology</i> , 2020, 8, 830.	1.8	14
9	Does SARS-CoV-2 has a longer incubation period than SARS and MERS?. <i>Journal of Medical Virology</i> , 2020, 92, 476-478.	2.5	203
10	WDR5 Facilitates Human Cytomegalovirus Replication by Promoting Capsid Nuclear Egress. <i>Journal of Virology</i> , 2018, 92, .	1.5	20
11	Serologic and viral genome prevalence of HSV, EBV, and HCMV among healthy adults in Wuhan, China. <i>Journal of Medical Virology</i> , 2018, 90, 571-581.	2.5	15
12	Tick-Borne Encephalitis Virus Nonstructural Protein NS5 Induces RANTES Expression Dependent on the RNA-Dependent RNA Polymerase Activity. <i>Journal of Immunology</i> , 2018, 201, 53-68.	0.4	30
13	Human Cytomegalovirus Immediate Early 1 Protein Causes Loss of SOX2 from Neural Progenitor Cells by Trapping Unphosphorylated STAT3 in the Nucleus. <i>Journal of Virology</i> , 2018, 92, .	1.5	20
14	Proteomic Analysis of Zika Virus Infected Primary Human Fetal Neural Progenitors Suggests a Role for Doublecortin in the Pathological Consequences of Infection in the Cortex. <i>Frontiers in Microbiology</i> , 2018, 9, 1067.	1.5	37
15	Human cytomegalovirus infection dysregulates neural progenitor cell fate by disrupting Hes1 rhythm and down-regulating its expression. <i>Virologica Sinica</i> , 2017, 32, 188-198.	1.2	9
16	Anterograde monosynaptic transneuronal tracers derived from herpes simplex virus 1 strain H129. <i>Molecular Neurodegeneration</i> , 2017, 12, 38.	4.4	94
17	ORF7 of Varicella-Zoster Virus Is Required for Viral Cytoplasmic Envelopment in Differentiated Neuronal Cells. <i>Journal of Virology</i> , 2017, 91, .	1.5	26
18	Infected T98G glioblastoma cells support human cytomegalovirus reactivation from latency. <i>Virology</i> , 2017, 510, 205-215.	1.1	8

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19	Human cytomegalovirus IE1 downregulates Hes1 in neural progenitor cells as a potential E3 ubiquitin ligase. <i>PLoS Pathogens</i> , 2017, 13, e1006542.	2.1	38
20	Transcriptional Activation of Multiple Operons Involved in para-Nitrophenol Degradation by <i>Pseudomonas</i> sp. Strain WBC-3. <i>Applied and Environmental Microbiology</i> , 2015, 81, 220-230.	1.4	31
21	Two Polyhydroxyalkanoate Synthases from Distinct Classes from the Aromatic Degradier <i>Cupriavidus pinatubonensis</i> JMP134 Exhibit the Same Substrate Preference. <i>PLoS ONE</i> , 2015, 10, e0142332.	1.1	5
22	Construction of an engineered strain capable of degrading two isomeric nitrophenols via a <i>sacB</i> - and <i>gfp</i> -based markerless integration system. <i>Applied Microbiology and Biotechnology</i> , 2014, 98, 4749-4756.	1.7	14
23	Intragenomic Heterogeneity of 16S rRNA Genes Causes Overestimation of Prokaryotic Diversity. <i>Applied and Environmental Microbiology</i> , 2013, 79, 5962-5969.	1.4	341