

Bo Liang

List of Publications by Year in descending order

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33
papers

1,669
citations

430754

18
h-index

434063

31
g-index

34
all docs

34
docs citations

34
times ranked

2140
citing authors

#	ARTICLE	IF	CITATIONS
1	Cryo-Electron Microscopy Structures of the <i>Pneumoviridae</i> Polymerases. <i>Viral Immunology</i> , 2021, 34, 18-26.	0.6	3
2	Efficient purification and assembly of ribonucleoprotein complex for interaction analysis by MST assay coupled with GaMD simulations. <i>STAR Protocols</i> , 2021, 2, 100315.	0.5	2
3	Baicalein and Baicalin Inhibit SARS-CoV-2 RNA-Dependent-RNA Polymerase. <i>Microorganisms</i> , 2021, 9, 893.	1.6	80
4	Structural Insights into the Respiratory Syncytial Virus RNA Synthesis Complexes. <i>Viruses</i> , 2021, 13, 834.	1.5	15
5	Generation and Assembly of Virus-Specific Nucleocapsids of the Respiratory Syncytial Virus. <i>Journal of Visualized Experiments</i> , 2021, , .	0.2	0
6	Discovery of the first chemical tools to regulate MKK3-mediated MYC activation in cancer. <i>Bioorganic and Medicinal Chemistry</i> , 2021, 45, 116324.	1.4	8
7	<i>In vitro</i> trackable assembly of RNA-specific nucleocapsids of the respiratory syncytial virus. <i>Journal of Biological Chemistry</i> , 2020, 295, 883-895.	1.6	9
8	Structure of the Vesicular Stomatitis Virus L Protein in Complex with Its Phosphoprotein Cofactor. <i>Cell Reports</i> , 2020, 30, 53-60.e5.	2.9	51
9	<i>In Vitro</i> Primer-Based RNA Elongation and Promoter Fine Mapping of the Respiratory Syncytial Virus. <i>Journal of Virology</i> , 2020, 95, .	1.5	9
10	Repurposing Nucleoside Analogs for Human Coronaviruses. <i>Antimicrobial Agents and Chemotherapy</i> , 2020, 65, .	1.4	45
11	Structure of the Human Respiratory Syncytial Virus M2-1 Protein in Complex with a Short Positive-Sense Gene-End RNA. <i>Structure</i> , 2020, 28, 979-990.e4.	1.6	15
12	Structures of the <i>Mononegavirales</i> Polymerases. <i>Journal of Virology</i> , 2020, 94, .	1.5	28
13	Cryo-EM structure of the respiratory syncytial virus RNA polymerase. <i>Nature Communications</i> , 2020, 11, 368.	5.8	61
14	<i>In vitro</i> trackable assembly of RNA-specific nucleocapsids of the respiratory syncytial virus. <i>Journal of Biological Chemistry</i> , 2020, 295, 883-895.	1.6	8
15	A conserved histidine in Group-1 influenza subtype hemagglutinin proteins is essential for membrane fusion activity. <i>Virology</i> , 2019, 536, 78-90.	1.1	11
16	Cryo-EM structure of TRPC5 at 2.8-Å... resolution reveals unique and conserved structural elements essential for channel function. <i>Science Advances</i> , 2019, 5, eaaw7935.	4.7	69
17	Activation of the Hemagglutinin of Influenza Viruses. , 2018, , 3-26.		3
18	An <i>In Vitro</i> RNA Synthesis Assay for Rabies Virus Defines Ribonucleoprotein Interactions Critical for Polymerase Activity. <i>Journal of Virology</i> , 2017, 91, .	1.5	30

#	ARTICLE	IF	CITATIONS
19	Peripheral motor neuropathy is associated with defective kinase regulation of the KCC3 cotransporter. <i>Science Signaling</i> , 2016, 9, ra77.	1.6	46
20	Regulatory domain or CpG site variation in SLC12A5, encoding the chloride transporter KCC2, in human autism and schizophrenia. <i>Frontiers in Cellular Neuroscience</i> , 2015, 9, 386.	1.8	86
21	Structure of the L Protein of Vesicular Stomatitis Virus from Electron Cryomicroscopy. <i>Cell</i> , 2015, 162, 314-327.	13.5	211
22	Genetically encoded impairment of neuronal KCC2 cotransporter function in human idiopathic generalized epilepsy. <i>EMBO Reports</i> , 2014, 15, 766-774.	2.0	163
23	Modulation of neuronal activity by phosphorylation of the Cl ⁻ cotransporter KCC2. <i>Trends in Neurosciences</i> , 2013, 36, 726-737.	4.2	196
24	Critical phosphoprotein elements that regulate polymerase architecture and function in vesicular stomatitis virus. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 14628-14633.	3.3	57
25	Structures of ribonucleoprotein particle modification enzymes. <i>Quarterly Reviews of Biophysics</i> , 2011, 44, 95-122.	2.4	20
26	Structural and functional evidence of high specificity of Cbf5 for ACA trinucleotide. <i>Rna</i> , 2011, 17, 244-250.	1.6	13
27	Molecular architecture of the vesicular stomatitis virus RNA polymerase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 20075-20080.	3.3	91
28	Functional and Structural Impact of Target Uridine Substitutions on the H/ACA Ribonucleoprotein Particle Pseudouridine Synthase. <i>Biochemistry</i> , 2010, 49, 6276-6281.	1.2	19
29	Glycosidic Bond Conformation Preference Plays a Pivotal Role in Catalysis of RNA Pseudouridylation: A Combined Simulation and Structural Study. <i>Journal of Molecular Biology</i> , 2010, 401, 690-695.	2.0	11
30	Structure of a functional ribonucleoprotein pseudouridine synthase bound to a substrate RNA. <i>Nature Structural and Molecular Biology</i> , 2009, 16, 740-746.	3.6	77
31	Long-distance placement of substrate RNA by H/ACA proteins. <i>Rna</i> , 2008, 14, 2086-2094.	1.6	22
32	Substrate RNA positioning in the archaeal H/ACA ribonucleoprotein complex. <i>Nature Structural and Molecular Biology</i> , 2007, 14, 1189-1195.	3.6	57
33	Crystal Structure of a Cbf5-Nop10-Gar1 Complex and Implications in RNA-Guided Pseudouridylation and Dyskeratosis Congenita. <i>Molecular Cell</i> , 2006, 21, 249-260.	4.5	152