Jianping Wu

List of Publications by Year in descending order

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279778 330122 4,312 43 23 37 h-index citations g-index papers 47 47 47 6183 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Crystal structure of the human glucose transporter GLUT1. Nature, 2014, 510, 121-125.	27.8	592
2	Structure of the voltage-gated calcium channel Cav1.1 at 3.6 à resolution. Nature, 2016, 537, 191-196.	27.8	398
3	Structure of the rabbit ryanodine receptor RyR1 at near-atomic resolution. Nature, 2015, 517, 50-55.	27.8	391
4	Structure of a eukaryotic voltage-gated sodium channel at near-atomic resolution. Science, 2017, 355, .	12.6	351
5	Structure of the Nav1.4-β1 Complex from Electric Eel. Cell, 2017, 170, 470-482.e11.	28.9	272
6	Structure of the voltage-gated calcium channel Ca $<$ sub $>$ v $<$ /sub $>$ 1.1 complex. Science, 2015, 350, aad2395.	12.6	270
7	Structural Insights into the Niemann-Pick C1 (NPC1)-Mediated Cholesterol Transfer and Ebola Infection. Cell, 2016, 165, 1467-1478.	28.9	266
8	Structural basis for the gating mechanism of the type 2 ryanodine receptor RyR2. Science, 2016, 354, .	12.6	221
9	Structural basis for the modulation of voltage-gated sodium channels by animal toxins. Science, 2018, 362, .	12.6	200
10	An atomic structure of the human 26S proteasome. Nature Structural and Molecular Biology, 2016, 23, 778-785.	8.2	189
11	Molecular Basis for Ligand Modulation of a Mammalian Voltage-Gated Ca2+ Channel. Cell, 2019, 177, 1495-1506.e12.	28.9	172
12	Functional and genetic analysis of viral receptor ACE2 orthologs reveals a broad potential host range of SARS-CoV-2. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	168
13	TIPE3 Is the Transfer Protein of Lipid Second Messengers that Promote Cancer. Cancer Cell, 2014, 26, 465-478.	16.8	93
14	The Central domain of RyR1 is the transducer for long-range allosteric gating of channel opening. Cell Research, 2016, 26, 995-1006.	12.0	93
15	Structure of an endogenous yeast 26S proteasome reveals two major conformational states. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 2642-2647.	7.1	75
16	Crystal structure of human lysyl oxidase-like 2 (hLOXL2) in a precursor state. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 3828-3833.	7.1	55
17	Structure of a mammalian sperm cation channel complex. Nature, 2021, 595, 746-750.	27.8	44
18	Structural basis for gating mechanism of Pannexin 1 channel. Cell Research, 2020, 30, 452-454.	12.0	43

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19	Mutation Y453F in the spike protein of SARS-CoV-2 enhances interaction with the mink ACE2 receptor for host adaption. PLoS Pathogens, 2021, 17, e1010053.	4.7	43
20	Structural and biochemical analysis of Bcl-2 interaction with the hepatitis B virus protein HBx. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 2074-2079.	7.1	42
21	Crystal structure of a LacY–nanobody complex in a periplasmic-open conformation. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 12420-12425.	7.1	38
22	Comparative analysis reveals the species-specific genetic determinants of ACE2 required for SARS-CoV-2 entry. PLoS Pathogens, 2021, 17, e1009392.	4.7	34
23	Crystal structure of a mycobacterial Insig homolog provides insight into how these sensors monitor sterol levels. Science, 2015, 349, 187-191.	12.6	32
24	Structure of the human sodium leak channel NALCN in complex with FAM155A. Nature Communications, 2020, 11, 5831.	12.8	24
25	Susceptibilities of Human ACE2 Genetic Variants in Coronavirus Infection. Journal of Virology, 2022, 96, JVI0149221.	3.4	22
26	Examination of the Dimerization States of the Single-stranded RNA Recognition Protein Pentatricopeptide Repeat 10 (PPR10). Journal of Biological Chemistry, 2014, 289, 31503-31512.	3.4	16
27	Cryo-EM structures of human TMEM120A and TMEM120B. Cell Discovery, 2021, 7, 77.	6.7	16
28	Structure of the WD40 domain of SCAP from fission yeast reveals the molecular basis for SREBP recognition. Cell Research, 2015, 25, 401-411.	12.0	15
29	Adaptive Traffic Signal Control Model on Intersections Based on Deep Reinforcement Learning. Journal of Advanced Transportation, 2020, 2020, 1-14.	1.7	11
30	Preventing IP source address spoofing: A two-level, state machine-based method. Tsinghua Science and Technology, 2009, 14, 413-422.	6.1	10
31	Structure-Function Relationship of the Voltage-Gated Calcium Channel Cav1.1 Complex. Advances in Experimental Medicine and Biology, 2017, 981, 23-39.	1.6	9
32	Factors Influencing the Acceptance of Robo-Taxi Services in China: An Extended Technology Acceptance Model Analysis. Journal of Advanced Transportation, 2022, 2022, 1-11.	1.7	8
33	A two-level source address spoofing prevention based on automatic signature and verification mechanism., 2008,,.		7
34	Architecture of the human NALCN channelosome. Cell Discovery, 2022, 8, 33.	6.7	7
35	A scalable, Web-based architecture for hierarchical network management. , 0, , .		6
36	A Study on Public Adoption of Robo-Taxis in China. Journal of Advanced Transportation, 2020, 2020, 1-8.	1.7	6

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37	High-fidelity biosensing of dNTPs and nucleic acids by controllable subnanometer channel PaMscS. Biosensors and Bioelectronics, 2022, 200, 113894.	10.1	6
38	Exploring the Energy Efficiency of Electric Vehicles with Driving Behavioral Data from a Field Test and Questionnaire. Journal of Advanced Transportation, 2018, 2018, 1-14.	1.7	5
39	IPv6 development in China., 0,,.		4
40	An IPv6 Test-Bed Implementation for a Future Source Address Validation Architecture., 2008,,.		3
41	Online Traffic Accident Spatial-Temporal Post-Impact Prediction Model on Highways Based on Spiking Neural Networks. Journal of Advanced Transportation, 2021, 2021, 1-20.	1.7	2
42	A Web-based, event-driven management architecture. , 1999, , .		1
43	CNGI Project and CERNET2., 2004, , .		1