Yan Zhao

List of Publications by Year in descending order

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		430754	477173
29	1,781	18	29
papers	citations	h-index	g-index
32	32	32	2835
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Structure of human glycosylphosphatidylinositol transamidase. Nature Structural and Molecular Biology, 2022, 29, 203-209.	3.6	11
2	Structural basis for modulation of human NaV1.3 by clinical drug and selective antagonist. Nature Communications, 2022, 13, 1286.	5.8	36
3	Structure, gating, and pharmacology of human CaV3.3 channel. Nature Communications, 2022, 13, 2084.	5.8	22
4	Structural basis of autoinhibition of the human NHE3-CHP1 complex. Science Advances, 2022, 8, .	4.7	11
5	Structural basis of ligand binding modes of human EAAT2. Nature Communications, 2022, 13, .	5.8	12
6	Structure and mechanism of the human NHE1-CHP1 complex. Nature Communications, 2021, 12, 3474.	5.8	45
7	SLC-30A9 is required for Zn ²⁺ homeostasis, Zn ²⁺ mobilization, and mitochondrial health. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	21
8	TMEM120A contains a specific coenzyme A-binding site and might not mediate poking- or stretch-induced channel activities in cells. ELife, 2021, 10, .	2.8	20
9	Kainate receptor modulation by NETO2. Nature, 2021, 599, 325-329.	13.7	20
10	Closed-state inactivation and pore-blocker modulation mechanisms of human CaV2.2. Cell Reports, 2021, 37, 109931.	2.9	35
11	Structure of the Cardiac Sodium Channel. Cell, 2020, 180, 122-134.e10.	13.5	217
12	Structure of theDietziaMrp complex reveals molecular mechanism of this giant bacterial sodium proton pump. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 31166-31176.	3.3	8
13	Architecture and subunit arrangement of native AMPA receptors elucidated by cryo-EM. Science, 2019, 364, 355-362.	6.0	134
14	Structure of YidC from <i>Thermotoga maritima</i> and its implications for YidCâ€mediated membrane protein insertion. FASEB Journal, 2018, 32, 2411-2421.	0.2	28
15	Etoposide-induced protein 2.4 functions as a regulator of the calcium ATPase and protects pancreatic β-cell survival. Journal of Biological Chemistry, 2018, 293, 10128-10140.	1.6	18
16	Activation and Desensitization Mechanism of AMPA Receptor-TARP Complex by Cryo-EM. Cell, 2017, 170, 1234-1246.e14.	13.5	111
17	Architecture of fully occupied GluA2 AMPA receptor–TARP complex elucidated by cryo-EM. Nature, 2016, 536, 108-111.	13.7	100
18	Crystal structure of E. coli lipoprotein diacylglyceryl transferase. Nature Communications, 2016, 7, 10198.	5.8	81

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19	Thermodynamics of ABC transporters. Protein and Cell, 2016, 7, 17-27.	4.8	19
20	Homeostatic Control of Innate Lung Inflammation by Vici Syndrome Gene Epg5 and Additional Autophagy Genes Promotes Influenza Pathogenesis. Cell Host and Microbe, 2016, 19, 102-113.	5.1	83
21	Energy coupling mechanisms of <scp>MFS</scp> transporters. Protein Science, 2015, 24, 1560-1579.	3.1	101
22	Substrate-bound structure of the E. coli multidrug resistance transporter MdfA. Cell Research, 2015, 25, 1060-1073.	5.7	149
23	Crystal structure of lipid phosphatase <i>Escherichia coli</i> phosphatidylglycerophosphate phosphatase B. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 7636-7640.	3.3	52
24	Dapper1 promotes autophagy by enhancing the Beclin1-Vps34-Atg14L complex formation. Cell Research, 2014, 24, 912-924.	5.7	57
25	Structure of the nonameric bacterial amyloid secretion channel. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, E5439-44.	3.3	87
26	Crystal structure and biochemical studies of Brucella melitensis 5′-methylthioadenosine/S-adenosylhomocysteine nucleosidase. Biochemical and Biophysical Research Communications, 2014, 446, 965-970.	1.0	6
27	Atomic resolution structure of the E. coli YajR transporter YAM domain. Biochemical and Biophysical Research Communications, 2014, 450, 929-935.	1.0	8
28	Crystal Structure of the E.Âcoli Peptide Transporter YbgH. Structure, 2014, 22, 1152-1160.	1.6	66
29	Structural basis for lipopolysaccharide insertion in the bacterial outer membrane. Nature, 2014, 511, 108-111	13.7	221