

# Baoyu Chen

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5442335/publications.pdf>

Version: 2024-02-01

28  
papers

3,484  
citations

516710

16  
h-index

552781

26  
g-index

34  
all docs

34  
docs citations

34  
times ranked

5035  
citing authors

#	ARTICLE	IF	CITATIONS
1	WASP family proteins: Molecular mechanisms and implications in human disease. <i>European Journal of Cell Biology</i> , 2022, 101, 151244.	3.6	19
2	WAVE regulatory complex. <i>Current Biology</i> , 2021, 31, R512-R517.	3.9	60
3	A two-step actin polymerization mechanism drives dendrite branching. <i>Neural Development</i> , 2021, 16, 3.	2.4	10
4	HEM1 deficiency disrupts mTORC2 and F-actin control in inherited immunodysregulatory disease. <i>Science</i> , 2020, 369, 202-207.	12.6	65
5	Endosomal receptor trafficking: Retromer and beyond. <i>Traffic</i> , 2018, 19, 578-590.	2.7	133
6	A Dendritic Guidance Receptor Complex Brings Together Distinct Actin Regulators to Drive Efficient F-Actin Assembly and Branching. <i>Developmental Cell</i> , 2018, 45, 362-375.e3.	7.0	56
7	Rac1 GTPase activates the WAVE regulatory complex through two distinct binding sites. <i>ELife</i> , 2017, 6, .	6.0	129
8	Fat2 acts through the WAVE regulatory complex to drive collective cell migration during tissue rotation. <i>Journal of Cell Biology</i> , 2016, 212, 591-603.	5.2	54
9	Biochemical Reconstitution of the WAVE Regulatory Complex. <i>Methods in Enzymology</i> , 2014, 540, 55-72.	1.0	20
10	Local F-actin Network Links Synapse Formation and Axon Branching. <i>Cell</i> , 2014, 156, 208-220.	28.9	128
11	The WAVE Regulatory Complex Links Diverse Receptors to the Actin Cytoskeleton. <i>Cell</i> , 2014, 156, 195-207.	28.9	260
12	Ena/VASP Proteins Cooperate with the WAVE Complex to Regulate the Actin Cytoskeleton. <i>Developmental Cell</i> , 2014, 30, 569-584.	7.0	101
13	Phase transitions in the assembly of multivalent signalling proteins. <i>Nature</i> , 2012, 483, 336-340.	27.8	1,938
14	Engagement of Arginine Finger to ATP Triggers Large Conformational Changes in NtrC1 AAA+ ATPase for Remodeling Bacterial RNA Polymerase. <i>Structure</i> , 2010, 18, 1420-1430.	3.3	49
15	The WAVE regulatory complex is inhibited. <i>Nature Structural and Molecular Biology</i> , 2009, 16, 561-563.	8.2	135
16	ADPase activity of recombinantly expressed thermotolerant ATPases may be caused by copurification of adenylate kinase of <i>Escherichia coli</i> . <i>FEBS Journal</i> , 2009, 276, 807-815.	4.7	8
17	Sequential Action of ATP on the Enhancer Binding AAA+ ATPase NtrC1. <i>FASEB Journal</i> , 2009, 23, 495.21.	0.5	0
18	Regulation and action of the bacterial enhancer-binding protein AAA+ domains. <i>Biochemical Society Transactions</i> , 2008, 36, 89-93.	3.4	16

#	ARTICLE	IF	CITATIONS
19	ATP Ground- and Transition States of Bacterial Enhancer Binding AAA+ ATPases Support Complex Formation with Their Target Protein, <i>Jf54</i> . <i>Structure</i> , 2007, 15, 429-440.	3.3	64
20	The structural basis for regulated assembly and function of the transcriptional activator NtrC. <i>Genes and Development</i> , 2006, 20, 1485-1495.	5.9	109
21	Molecular mechanisms of hormonal activity. I. receptors. neuromediators. systems with second messengers. <i>Biochemistry (Moscow)</i> , 2005, 70, 24-39.	1.5	0
22	Negative Regulation of AAA+ ATPase Assembly by Two Component Receiver Domains: A Transcription Activation Mechanism that is Conserved in Mesophilic and Extremely Hyperthermophilic Bacteria. <i>Journal of Molecular Biology</i> , 2005, 353, 242-255.	4.2	53
23	Evidence for proximal cysteine and lysine residues at or near the active site of arginine kinase of <i>Stichopus japonicus</i> . <i>Biochemistry (Moscow)</i> , 2004, 69, 1336-1343.	1.5	10
24	Urea Induced Inactivation and Unfolding of Arginine Kinase from the Sea Cucumber <i>Stichopus japonicus</i> . <i>Biochemistry (Moscow)</i> , 2003, 68, 1267-1271.	1.5	5
25	Multiple effects of chemical reagent on enzyme: o-phthalaldehyde-induced inactivation, dissociation and partial unfolding of lactate dehydrogenase from pig heart. <i>International Journal of Biological Macromolecules</i> , 2003, 32, 191-197.	7.5	8
26	Expression, purification, and characterization of arginine kinase from the sea cucumber <i>Stichopus japonicus</i> . <i>Protein Expression and Purification</i> , 2003, 29, 230-234.	1.3	24
27	Inactivation and conformational changes of lactate dehydrogenase from porcine heart in sodium dodecyl sulfate solutions. <i>International Journal of Biological Macromolecules</i> , 2002, 31, 97-102.	7.5	9
28	p-Chloromercuribenzoate-induced inactivation and partial unfolding of porcine heart lactate dehydrogenase. <i>Biochemistry (Moscow)</i> , 2002, 67, 583-587.	1.5	3