## John S Allingham

## List of Publications by Citations

Source: https://exaly.com/author-pdf/2834694/john-s-allingham-publications-by-citations.pdf

Version: 2024-04-17

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

31	<b>1,181</b> citations	17	<b>32</b>
papers		h-index	g-index
32 ext. papers	1,350 ext. citations	10.1 avg, IF	4.03 L-index

#	Paper	IF	Citations
31	The structural basis of blebbistatin inhibition and specificity for myosin II. <i>Nature Structural and Molecular Biology</i> , <b>2005</b> , 12, 378-9	17.6	228
30	Trisoxazole macrolide toxins mimic the binding of actin-capping proteins to actin. <i>Nature Structural and Molecular Biology</i> , <b>2003</b> , 10, 1058-63	17.6	134
29	An antifreeze protein folds with an interior network of more than 400 semi-clathrate waters. <i>Science</i> , <b>2014</b> , 343, 795-8	33.3	122
28	Structures of microfilament destabilizing toxins bound to actin provide insight into toxin design and activity. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2005</b> , 102, 14527-32	11.5	8o
27	Vik1 modulates microtubule-Kar3 interactions through a motor domain that lacks an active site. <i>Cell</i> , <b>2007</b> , 128, 1161-72	56.2	72
26	The RAG1/RAG2 complex constitutes a 3Vflap endonuclease: implications for junctional diversity in V(D)J and transpositional recombination. <i>Molecular Cell</i> , <b>1999</b> , 4, 935-47	17.6	70
25	A structural basis for regulation of actin polymerization by pectenotoxins. <i>Journal of Molecular Biology</i> , <b>2007</b> , 371, 959-70	6.5	58
24	Structure of a 1.5-MDa adhesin that binds its Antarctic bacterium to diatoms and ice. <i>Science Advances</i> , <b>2017</b> , 3, e1701440	14.3	52
23	Absolute stereochemistry of ulapualide A. <i>Organic Letters</i> , <b>2004</b> , 6, 597-9	6.2	45
23	Absolute stereochemistry of ulapualide A. <i>Organic Letters</i> , <b>2004</b> , 6, 597-9  All three residues of the Tn 10 transposase DDE catalytic triad function in divalent metal ion binding. <i>Journal of Molecular Biology</i> , <b>1999</b> , 289, 1195-206	6.2	45 45
	All three residues of the Tn 10 transposase DDE catalytic triad function in divalent metal ion		45
22	All three residues of the Tn 10 transposase DDE catalytic triad function in divalent metal ion binding. <i>Journal of Molecular Biology</i> , <b>1999</b> , 289, 1195-206  Actin-binding cleft closure in myosin II probed by site-directed spin labeling and pulsed EPR.	6.5	45
22	All three residues of the Tn 10 transposase DDE catalytic triad function in divalent metal ion binding. <i>Journal of Molecular Biology</i> , <b>1999</b> , 289, 1195-206  Actin-binding cleft closure in myosin II probed by site-directed spin labeling and pulsed EPR. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2008</b> , 105, 12867-72  KIF14 binds tightly to microtubules and adopts a rigor-like conformation. <i>Journal of Molecular</i>	6.5	45 41
22 21 20	All three residues of the Tn 10 transposase DDE catalytic triad function in divalent metal ion binding. <i>Journal of Molecular Biology</i> , <b>1999</b> , 289, 1195-206  Actin-binding cleft closure in myosin II probed by site-directed spin labeling and pulsed EPR. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2008</b> , 105, 12867-72  KIF14 binds tightly to microtubules and adopts a rigor-like conformation. <i>Journal of Molecular Biology</i> , <b>2014</b> , 426, 2997-3015  The small molecule tool (S)-(-)-blebbistatin: novel insights of relevance to myosin inhibitor design.	6.5 11.5 6.5	45 41 33 29
22 21 20	All three residues of the Tn 10 transposase DDE catalytic triad function in divalent metal ion binding. <i>Journal of Molecular Biology</i> , <b>1999</b> , 289, 1195-206  Actin-binding cleft closure in myosin II probed by site-directed spin labeling and pulsed EPR. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2008</b> , 105, 12867-72  KIF14 binds tightly to microtubules and adopts a rigor-like conformation. <i>Journal of Molecular Biology</i> , <b>2014</b> , 426, 2997-3015  The small molecule tool (S)-(-)-blebbistatin: novel insights of relevance to myosin inhibitor design. <i>Organic and Biomolecular Chemistry</i> , <b>2008</b> , 6, 2076-84  Ternary complex of Kif2A-bound tandem tubulin heterodimers represents a kinesin-13-mediated	6.5 11.5 6.5 3.9	45 41 33 29
22 21 20 19	All three residues of the Tn 10 transposase DDE catalytic triad function in divalent metal ion binding. <i>Journal of Molecular Biology</i> , <b>1999</b> , 289, 1195-206  Actin-binding cleft closure in myosin II probed by site-directed spin labeling and pulsed EPR. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2008</b> , 105, 12867-72  KIF14 binds tightly to microtubules and adopts a rigor-like conformation. <i>Journal of Molecular Biology</i> , <b>2014</b> , 426, 2997-3015  The small molecule tool (S)-(-)-blebbistatin: novel insights of relevance to myosin inhibitor design. <i>Organic and Biomolecular Chemistry</i> , <b>2008</b> , 6, 2076-84  Ternary complex of Kif2A-bound tandem tubulin heterodimers represents a kinesin-13-mediated microtubule depolymerization reaction intermediate. <i>Nature Communications</i> , <b>2018</b> , 9, 2628	6.5 11.5 6.5 3.9	45 41 33 29 27

## LIST OF PUBLICATIONS

14	Role of Ca⊞ in folding the tandem □sandwich extender domains of a bacterial ice-binding adhesin. <i>FEBS Journal</i> , <b>2013</b> , 280, 5919-32	5.7	17	
13	Conditional switching of KIF2A mutation provides new insights into cortical malformation pathogeny. <i>Human Molecular Genetics</i> , <b>2020</b> , 29, 766-784	5.6	9	
12	Neck rotation and neck mimic docking in the noncatalytic Kar3-associated protein Vik1. <i>Journal of Biological Chemistry</i> , <b>2012</b> , 287, 40292-301	5.4	9	
11	Peptide backbone circularization enhances antifreeze protein thermostability. <i>Protein Science</i> , <b>2017</b> , 26, 1932-1941	6.3	7	
10	Crystal structure of the Kar3-like kinesin motor domain from the filamentous fungus Ashbya gossypii. <i>Proteins: Structure, Function and Bioinformatics</i> , <b>2012</b> , 80, 1016-27	4.2	7	
9	These motors were made for walking. <i>Protein Science</i> , <b>2020</b> , 29, 1707-1723	6.3	5	
8	Functional adaptation between yeast actin and its cognate myosin motors. <i>Journal of Biological Chemistry</i> , <b>2011</b> , 286, 30384-30392	5.4	5	
7	Crystal structure of the Candida albicans Kar3 kinesin motor domain fused to maltose-binding protein. <i>Biochemical and Biophysical Research Communications</i> , <b>2012</b> , 428, 427-32	3.4	4	
6	Kar3Vik1 mechanochemistry is inhibited by mutation or deletion of the C terminus of the Vik1 subunit. <i>Journal of Biological Chemistry</i> , <b>2013</b> , 288, 36957-70	5.4	4	
5	Candida albicans Kinesin Kar3 Depends on a Cik1-Like Regulatory Partner Protein for Its Roles in Mating, Cell Morphogenesis, and Bipolar Spindle Formation. <i>Eukaryotic Cell</i> , <b>2015</b> , 14, 755-74		3	
4	Actin-binding toxin "tail" wags the dog. <i>Chemistry and Biology</i> , <b>2008</b> , 15, 205-7		3	
3	Phasing with calcium at home. <i>Acta Crystallographica Section F, Structural Biology Communications</i> , <b>2019</b> , 75, 377-384	1.1	2	
2	Ste2 receptor-mediated chemotropism of Fusarium graminearum contributes to its pathogenicity against wheat. <i>Scientific Reports</i> , <b>2020</b> , 10, 10770	4.9	2	
1	Kinesin-5 Is Dispensable for Bipolar Spindle Formation and Elongation in Candida albicans, but Simultaneous Loss of Kinesin-14 Activity Is Lethal. <i>MSphere</i> , <b>2019</b> , 4,	5	1	