

Dimitrios Vavylonis

List of Publications by Citations

Source: <https://exaly.com/author-pdf/8043928/dimitrios-vavylonis-publications-by-citations.pdf>
Version: 2024-04-09

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.

64 papers	2,098 citations	24 h-index	45 g-index
83 ext. papers	2,646 ext. citations	5.9 avg, IF	5 L-index

#	Paper	IF	Citations
64	Assembly mechanism of the contractile ring for cytokinesis by fission yeast. <i>Science</i> , 2008 , 319, 97-100	33.3	294
63	Polymerization kinetics of ADP- and ADP-Pi-actin determined by fluorescence microscopy. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007 , 104, 8827-32	11.5	150
62	Model of formin-associated actin filament elongation. <i>Molecular Cell</i> , 2006 , 21, 455-66	17.6	144
61	Segmentation and tracking of cytoskeletal filaments using open active contours. <i>Cytoskeleton</i> , 2010 , 67, 693-705	2.4	124
60	Oscillatory dynamics of Cdc42 GTPase in the control of polarized growth. <i>Science</i> , 2012 , 337, 239-43	33.3	119
59	Actin polymerization kinetics, cap structure, and fluctuations. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005 , 102, 8543-8	11.5	105
58	Spontaneous Cdc42 polarization independent of GDI-mediated extraction and actin-based trafficking. <i>PLoS Biology</i> , 2015 , 13, e1002097	9.7	85
57	Nanoscale movements of cellulose microfibrils in primary cell walls. <i>Nature Plants</i> , 2017 , 3, 17056	11.5	79
56	Interactive, computer-assisted tracking of speckle trajectories in fluorescence microscopy: application to actin polymerization and membrane fusion. <i>Biophysical Journal</i> , 2011 , 101, 1794-804	2.9	68
55	Actinin and fimbrin cooperate with myosin II to organize actomyosin bundles during contractile-ring assembly. <i>Molecular Biology of the Cell</i> , 2012 , 23, 3094-110	3.5	67
54	Excitable actin dynamics in lamellipodial protrusion and retraction. <i>Biophysical Journal</i> , 2012 , 102, 1493-502	3.2	65
53	Reactive Polymer Interfaces: How Reaction Kinetics Depend on Reactivity and Density of Chemical Groups. <i>Macromolecules</i> , 1999 , 32, 1785-1796	5.5	56
52	SOAX: a software for quantification of 3D biopolymer networks. <i>Scientific Reports</i> , 2015 , 5, 9081	4.9	50
51	Two functionally distinct sources of actin monomers supply the leading edge of lamellipodia. <i>Cell Reports</i> , 2015 , 11, 433-45	10.6	43
50	Disentangling loosening from softening: insights into primary cell wall structure. <i>Plant Journal</i> , 2019 , 100, 1101-1117	6.9	42
49	A review of models of fluctuating protrusion and retraction patterns at the leading edge of motile cells. <i>Cytoskeleton</i> , 2012 , 69, 195-206	2.4	42
48	Local Pheromone Release from Dynamic Polarity Sites Underlies Cell-Cell Pairing during Yeast Mating. <i>Current Biology</i> , 2016 , 26, 1117-25	6.3	35

47	3D actin network centerline extraction with multiple active contours. <i>Medical Image Analysis</i> , 2014 , 18, 272-84	15.4	35
46	Dynamic network morphology and tension buildup in a 3D model of cytokinetic ring assembly. <i>Biophysical Journal</i> , 2014 , 107, 2618-28	2.9	35
45	New single-molecule speckle microscopy reveals modification of the retrograde actin flow by focal adhesions at nanometer scales. <i>Molecular Biology of the Cell</i> , 2014 , 25, 1010-24	3.5	32
44	ACTIN-DIRECTED TOXIN. ACD toxin-produced actin oligomers poison formin-controlled actin polymerization. <i>Science</i> , 2015 , 349, 535-9	33.3	31
43	Model of fission yeast cell shape driven by membrane-bound growth factors and the cytoskeleton. <i>PLoS Computational Biology</i> , 2013 , 9, e1003287	5	28
42	Distributed actin turnover in the lamellipodium and FRAP kinetics. <i>Biophysical Journal</i> , 2013 , 104, 247-57	2.9	27
41	AUTOMATED ACTIN FILAMENT SEGMENTATION, TRACKING AND TIP ELONGATION MEASUREMENTS BASED ON OPEN ACTIVE CONTOUR MODELS 2009 , 2009, 1302-1305	1.5	24
40	Model of For3p-mediated actin cable assembly in fission yeast. <i>PLoS ONE</i> , 2008 , 3, e4078	3.7	20
39	Interfacial reactions: mixed order kinetics and segregation effects. <i>Physical Review Letters</i> , 2000 , 84, 3193-6	7.4	19
38	Reconstitution of contractile actomyosin rings in vesicles. <i>Nature Communications</i> , 2021 , 12, 2254	17.4	19
37	Actin cable distribution and dynamics arising from cross-linking, motor pulling, and filament turnover. <i>Molecular Biology of the Cell</i> , 2014 , 25, 3006-16	3.5	18
36	Model of myosin node aggregation into a contractile ring: the effect of local alignment. <i>Journal of Physics Condensed Matter</i> , 2011 , 23, 374103	1.8	18
35	ER-PM Contacts Define Actomyosin Kinetics for Proper Contractile Ring Assembly. <i>Current Biology</i> , 2016 , 26, 647-53	6.3	15
34	Cytoskeletal dynamics in fission yeast: a review of models for polarization and division. <i>HFSP Journal</i> , 2010 , 4, 122-30		15
33	Myosin-dependent actin stabilization as revealed by single-molecule imaging of actin turnover. <i>Molecular Biology of the Cell</i> , 2018 , 29, 1941-1947	3.5	14
32	EXTRACTION AND ANALYSIS OF ACTIN NETWORKS BASED ON OPEN ACTIVE CONTOUR MODELS 2011 , 2011, 1334-1340	1.5	13
31	The ultrasensitivity of living polymers. <i>Physical Review Letters</i> , 2003 , 90, 118301	7.4	12
30	Actin Cross-Linking Toxin Is a Universal Inhibitor of Tandem-Organized and Oligomeric G-Actin Binding Proteins. <i>Current Biology</i> , 2018 , 28, 1536-1547.e9	6.3	11

29	Cell protrusion and retraction driven by fluctuations in actin polymerization: A two-dimensional model. <i>Cytoskeleton</i> , 2017 , 74, 490-503	2.4	11
28	Computational modeling highlights the role of the disordered Formin Homology 1 domain in profilin-actin transfer. <i>FEBS Letters</i> , 2018 , 592, 1804-1816	3.8	10
27	Kinetics of myosin node aggregation into a contractile ring. <i>Physical Review Letters</i> , 2010 , 105, 048102	7.4	10
26	Actin filament tracking based on particle filters and stretching open active contour models. <i>Lecture Notes in Computer Science</i> , 2009 , 12, 673-81	0.9	10
25	Exploration and stabilization of Ras1 mating zone: A mechanism with positive and negative feedbacks. <i>PLoS Computational Biology</i> , 2018 , 14, e1006317	5	10
24	Image analysis tools to quantify cell shape and protein dynamics near the leading edge. <i>Cell Structure and Function</i> , 2013 , 38, 1-7	2.2	9
23	Computational model of polarized actin cables and cytokinetic actin ring formation in budding yeast. <i>Cytoskeleton</i> , 2015 , 72, 517-33	2.4	8
22	Building a dendritic actin filament network branch by branch: models of filament orientation pattern and force generation in lamellipodia. <i>Biophysical Reviews</i> , 2018 , 10, 1577-1585	3.7	8
21	Organization of associating or crosslinked actin filaments in confinement. <i>Cytoskeleton</i> , 2019 , 76, 532-548	4.4	7
20	Convection-Induced Biased Distribution of Actin Probes in Live Cells. <i>Biophysical Journal</i> , 2019 , 116, 142-150	1.5	7
19	Molecular viewing of actin polymerizing actions and beyond: combination analysis of single-molecule speckle microscopy with modeling, FRAP and s-FDAP (sequential fluorescence decay after photoactivation). <i>Development Growth and Differentiation</i> , 2013 , 55, 508-14	3	6
18	Pulsed Laser Polymerization at Low Conversions: Broadening and Chain Transfer Effects. <i>Macromolecular Theory and Simulations</i> , 2003 , 12, 401-412	1.5	6
17	Model of turnover kinetics in the lamellipodium: implications of slow- and fast- diffusing capping protein and Arp2/3 complex. <i>Physical Biology</i> , 2016 , 13, 066009	3	6
16	Reconstitution of contractile actomyosin rings in vesicles		5
15	Lamellipodium tip actin barbed ends serve as a force sensor. <i>Genes To Cells</i> , 2019 , 24, 705-718	2.3	4
14	Formation of contractile networks and fibers in the medial cell cortex through myosin-II turnover, contraction, and stress-stabilization. <i>Cytoskeleton</i> , 2015 , 72, 29-46	2.4	4
13	Fission Yeast Polarization: Modeling Cdc42 Oscillations, Symmetry Breaking, and Zones of Activation and Inhibition. <i>Cells</i> , 2020 , 9,	7.9	4
12	Cell patterning by secretion-induced plasma membrane flows. <i>Science Advances</i> , 2021 , 7, eabg6718	14.3	4

11	Insights into Actin Polymerization and Nucleation Using a Coarse-Grained Model. <i>Biophysical Journal</i> , 2020 , 119, 553-566	2.9	3
10	Cell Biology: Capturing Formin's Mechano-Inhibition. <i>Current Biology</i> , 2017 , 27, R1078-R1080	6.3	2
9	Automated Tracking of Biopolymer Growth and Network Deformation with TSOAX. <i>Scientific Reports</i> , 2019 , 9, 1717	4.9	2
8	A systems-biology approach to yeast actin cables. <i>Advances in Experimental Medicine and Biology</i> , 2012 , 736, 325-35	3.6	2
7	Discrete mechanical model of lamellipodial actin network implements molecular clutch mechanism and generates arcs and microspikes. <i>PLoS Computational Biology</i> , 2021 , 17, e1009506	5	1
6	Cdc42 GTPase Activating Proteins (GAPs) Maintain Generational Inheritance of Cell Polarity and Cell Shape in Fission Yeast		1
5	Cell patterning by secretion-induced plasma membrane flows		1
4	Cdc42 GTPase-activating proteins (GAPs) regulate generational inheritance of cell polarity and cell shape in fission yeast. <i>Molecular Biology of the Cell</i> , 2021 , 32, ar14	3.5	1
3	Actin biophysics in the tradition of Fumio Oosawa: A special issue with contributions from participants at the 2016 "Now in Actin" meeting in Nagoya. <i>Cytoskeleton</i> , 2017 , 74, 445	2.4	
2	Molecular basis of cytokinesis in fission yeast. <i>FASEB Journal</i> , 2008 , 22, 115.2	0.9	
1	Lamellipodia in Stationary and Fluctuating States. <i>Modeling and Simulation in Science, Engineering and Technology</i> , 2018 , 211-258	0.8	