

# Hans G Othmer

## List of Publications by Year in descending order

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80  
papers

6,054  
citations

126858

33  
h-index

71651

76  
g-index

85  
all docs

85  
docs citations

85  
times ranked

4081  
citing authors

#	ARTICLE	IF	CITATIONS
1	The topology of the regulatory interactions predicts the expression pattern of the segment polarity genes in <i>Drosophila melanogaster</i> . <i>Journal of Theoretical Biology</i> , 2003, 223, 1-18.	0.8	827
2	Aggregation, Blowup, and Collapse: The ABC's of Taxis in Reinforced Random Walks. <i>SIAM Journal on Applied Mathematics</i> , 1997, 57, 1044-1081.	0.8	589
3	The Diffusion Limit of Transport Equations Derived from Velocity-Jump Processes. <i>SIAM Journal on Applied Mathematics</i> , 2000, 61, 751-775.	0.8	331
4	The Diffusion Limit of Transport Equations II: Chemotaxis Equations. <i>SIAM Journal on Applied Mathematics</i> , 2002, 62, 1222-1250.	0.8	297
5	Facilitated Transport of a Dpp/Scw Heterodimer by Sog/Tsg Leads to Robust Patterning of the <i>Drosophila</i> Blastoderm Embryo. <i>Cell</i> , 2005, 120, 873-886.	13.5	287
6	Mathematical modeling of tumor-induced angiogenesis. <i>Journal of Mathematical Biology</i> , 2004, 49, 111-87.	0.8	277
7	Shaping BMP morphogen gradients in the <i>Drosophila</i> embryo and pupal wing. <i>Development (Cambridge)</i> , 2006, 133, 183-193.	1.2	266
8	From Individual to Collective Behavior in Bacterial Chemotaxis. <i>SIAM Journal on Applied Mathematics</i> , 2004, 65, 361-391.	0.8	229
9	The BMP-Binding Protein Crossveinless 2 Is a Short-Range, Concentration-Dependent, Biphasic Modulator of BMP Signaling in <i>Drosophila</i> . <i>Developmental Cell</i> , 2008, 14, 940-953.	3.1	157
10	A HYBRID MODEL FOR TUMOR SPHEROID GROWTH <i>IN VITRO</i> I: THEORETICAL DEVELOPMENT AND EARLY RESULTS. <i>Mathematical Models and Methods in Applied Sciences</i> , 2007, 17, 1773-1798.	1.7	152
11	Multiscale Models of Taxis-Driven Patterning in Bacterial Populations. <i>SIAM Journal on Applied Mathematics</i> , 2009, 70, 133-167.	0.8	145
12	The role of the microenvironment in tumor growth and invasion. <i>Progress in Biophysics and Molecular Biology</i> , 2011, 106, 353-379.	1.4	145
13	A stochastic analysis of first-order reaction networks. <i>Bulletin of Mathematical Biology</i> , 2005, 67, 901-946.	0.9	144
14	A continuum model of motility in ameboid cells. <i>Bulletin of Mathematical Biology</i> , 2004, 66, 167-193.	0.9	132
15	A Mathematical Model for Outgrowth and Spatial Patterning of the Vertebrate Limb Bud. <i>Journal of Theoretical Biology</i> , 1999, 197, 295-330.	0.8	117
16	Robust, bistable patterning of the dorsal surface of the <i>Drosophila</i> embryo. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 11613-11618.	3.3	114
17	A discrete cell model with adaptive signalling for aggregation of <i>Dictyostelium discoideum</i> . <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 1997, 352, 391-417.	1.8	111
18	How cellular movement determines the collective force generated by the <i>Dictyostelium discoideum</i> slug. <i>Journal of Theoretical Biology</i> , 2004, 231, 203-222.	0.8	106

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19	From Signal Transduction to Spatial Pattern Formation in <i>E. coli</i> : A Paradigm for Multiscale Modeling in Biology. <i>Multiscale Modeling and Simulation</i> , 2005, 3, 362-394.	0.6	105
20	Mechanisms of scaling in pattern formation. <i>Development (Cambridge)</i> , 2013, 140, 4830-4843.	1.2	94
21	Multi-scale models of cell and tissue dynamics. <i>Philosophical Transactions Series A, Mathematical, Physical, and Engineering Sciences</i> , 2009, 367, 3525-3553.	1.6	93
22	Organism-Scale Modeling of Early <i>Drosophila</i> Patterning via Bone Morphogenetic Proteins. <i>Developmental Cell</i> , 2010, 18, 260-274.	3.1	85
23	A G protein-based model of adaptation in <i>Dictyostelium discoideum</i> . <i>Mathematical Biosciences</i> , 1994, 120, 25-76.	0.9	65
24	A Hybrid Model of Tumor-Stromal Interactions in Breast Cancer. <i>Bulletin of Mathematical Biology</i> , 2013, 75, 1304-1350.	0.9	62
25	The Intersection of Theory and Application in Elucidating Pattern Formation in Developmental Biology. <i>Mathematical Modelling of Natural Phenomena</i> , 2009, 4, 3-82.	0.9	61
26	Short- and long-range effects of Sonic hedgehog in limb development. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003, 100, 10152-10157.	3.3	54
27	Taxis equations for amoeboid cells. <i>Journal of Mathematical Biology</i> , 2007, 54, 847-885.	0.8	52
28	A multi-time-scale analysis of chemical reaction networks: I. Deterministic systems. <i>Journal of Mathematical Biology</i> , 2010, 60, 387-450.	0.8	51
29	The role of the tumor microenvironment in glioblastoma: A mathematical model. <i>IEEE Transactions on Biomedical Engineering</i> , 2016, 64, 1-1.	2.5	46
30	The effects of cell density and metabolite flux on cellular dynamics. <i>Journal of Mathematical Biology</i> , 1978, 5, 169-200.	0.8	42
31	A Theoretical Approach to Actin Filament Dynamics. <i>Journal of Statistical Physics</i> , 2007, 128, 111-138.	0.5	41
32	Robustness of Embryonic Spatial Patterning in <i>Drosophila melanogaster</i> . <i>Current Topics in Developmental Biology</i> , 2008, 81, 65-111.	1.0	41
33	Progress and perspectives in signal transduction, actin dynamics, and movement at the cell and tissue level: lessons from <i>Dictyostelium</i> . <i>Interface Focus</i> , 2016, 6, 20160047.	1.5	41
34	An equation-free computational approach for extracting population-level behavior from individual-based models of biological dispersal. <i>Physica D: Nonlinear Phenomena</i> , 2006, 215, 1-24.	1.3	35
35	Radial and Spiral Stream Formation in <i>Proteus mirabilis</i> Colonies. <i>PLoS Computational Biology</i> , 2011, 7, e1002332.	1.5	34
36	A Continuum Model of Actin Waves in <i>Dictyostelium discoideum</i> . <i>PLoS ONE</i> , 2013, 8, e64272.	1.1	34

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37	A mathematical model of salt-sensitive hypertension: the neurogenic hypothesis. <i>Journal of Physiology</i> , 2015, 593, 3065-3075.	1.3	32
38	Aggregation under local reinforcement: From lattice to continuum. <i>European Journal of Applied Mathematics</i> , 2004, 15, 545-576.	1.4	31
39	A Continuum Analysis of the Chemotactic Signal Seen by <i>Dictyostelium discoideum</i> . <i>Journal of Theoretical Biology</i> , 1998, 194, 461-483.	0.8	29
40	The Role of Mathematical Models in Understanding Pattern Formation in Developmental Biology. <i>Bulletin of Mathematical Biology</i> , 2015, 77, 817-845.	0.9	27
41	Hybrid models of cell and tissue dynamics in tumor growth. <i>Mathematical Biosciences and Engineering</i> , 2015, 12, 1141-1156.	1.0	25
42	Dynamic Receptor Team Formation Can Explain the High Signal Transduction Gain in <i>Escherichia coli</i> . <i>Biophysical Journal</i> , 2004, 86, 2650-2659.	0.2	23
43	A new method for choosing the computational cell in stochastic reaction-diffusion systems. <i>Journal of Mathematical Biology</i> , 2012, 65, 1017-1099.	0.8	23
44	A multi-time-scale analysis of chemical reaction networks: II. Stochastic systems. <i>Journal of Mathematical Biology</i> , 2016, 73, 1081-1129.	0.8	22
45	Excitation and Adaptation in Bacteria—a Model Signal Transduction System that Controls Taxis and Spatial Pattern Formation. <i>International Journal of Molecular Sciences</i> , 2013, 14, 9205-9248.	1.8	21
46	The discrete-time Kermack-McKendrick model: A versatile and computationally attractive framework for modeling epidemics. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	21
47	Computational analysis of amoeboid swimming at low Reynolds number. <i>Journal of Mathematical Biology</i> , 2016, 72, 1893-1926.	0.8	20
48	A Model for Direction Sensing in <i>Dictyostelium discoideum</i> : Ras Activity and Symmetry Breaking Driven by a $G_{i2}^{13}$ -Mediated, $G_{i2}$ -Rac8 -- Dependent Signal Transduction Network. <i>PLoS Computational Biology</i> , 2016, 12, e1004900.	1.5	20
49	A model for signal-relay adaptation in <i>Dictyostelium discoideum</i> . II. Analytical and numerical results. <i>Mathematical Biosciences</i> , 1985, 77, 79-139.	0.9	18
50	A stochastic analysis of actin polymerization in the presence of twinfilin and gelsolin. <i>Journal of Theoretical Biology</i> , 2007, 249, 723-736.	0.8	18
51	A model for pattern formation in <i>Dictyostelium discoideum</i> . <i>Differentiation</i> , 1996, 60, 1-16.	1.0	17
52	A Trimer of Dimers-Based Model for the Chemotactic Signal Transduction Network in Bacterial Chemotaxis. <i>Bulletin of Mathematical Biology</i> , 2012, 74, 2339-2382.	0.9	17
53	The Roles of Signaling in Cytoskeletal Changes, Random Movement, Direction-Sensing and Polarization of Eukaryotic Cells. <i>Cells</i> , 2020, 9, 1437.	1.8	16
54	The importance of geometry in mathematical models of developing systems. <i>Current Opinion in Genetics and Development</i> , 2012, 22, 547-552.	1.5	15

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55	Stochastic Analysis of Reaction-Diffusion Processes. Bulletin of Mathematical Biology, 2014, 76, 854-894.	0.9	15
56	Synergistic Effects of Bortezomib-OV Therapy and Anti-Invasive Strategies in Glioblastoma: A Mathematical Model. Cancers, 2019, 11, 215.	1.7	15
57	The effect of the signalling scheme on the robustness of pattern formation in development. Interface Focus, 2012, 2, 465-486.	1.5	14
58	Growth control in the Drosophila wing disk. Wiley Interdisciplinary Reviews: Systems Biology and Medicine, 2020, 12, e1478.	6.6	14
59	A theoretical analysis of filament length fluctuations in actin and other polymers. Journal of Mathematical Biology, 2011, 63, 1001-1049.	0.8	13
60	Getting in shape and swimming: the role of cortical forces and membrane heterogeneity in eukaryotic cells. Journal of Mathematical Biology, 2018, 77, 595-626.	0.8	13
61	Visualizing mesoderm and neural crest cell dynamics during chick head morphogenesis. Developmental Biology, 2020, 461, 184-196.	0.9	12
62	The Mathematical Analysis of Biological Aggregation and Dispersal: Progress, Problems and Perspectives. Lecture Notes in Mathematics, 2013, , 79-127.	0.1	11
63	Noise-induced mixing and multimodality in reaction networks. European Journal of Applied Mathematics, 2019, 30, 887-911.	1.4	11
64	Improving Parameter Inference from FRAP Data: an Analysis Motivated by Pattern Formation in the Drosophila Wing Disc. Bulletin of Mathematical Biology, 2017, 79, 448-497.	0.9	10
65	A Model for the Hippo Pathway in the Drosophila Wing Disc. Biophysical Journal, 2018, 115, 737-747.	0.2	10
66	Analysis of a model microswimmer with applications to blebbing cells and mini-robots. Journal of Mathematical Biology, 2018, 76, 1699-1763.	0.8	9
67	The performance of discrete models of low reynolds number swimmers. Mathematical Biosciences and Engineering, 2015, 12, 1303-1320.	1.0	8
68	Constant-complexity stochastic simulation algorithm with optimal binning. Journal of Chemical Physics, 2015, 143, 074108.	1.2	7
69	From Individual to Collective Behavior of Unicellular Organisms: Recent Results and Open Problems. , 2009, , .		5
70	Editorial: Special Issue on Stochastic Modelling of Reaction-Diffusion Processes in Biology. Bulletin of Mathematical Biology, 2014, 76, 761-765.	0.9	4
71	Scale invariance of morphogen-mediated patterning by flux optimization. , 2012, , .		3
72	A phosphoinositide-based model of actin waves in frustrated phagocytosis. Journal of Theoretical Biology, 2021, 527, 110764.	0.8	3

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73	How surrogates for cortical forces determine cell shape. <i>International Journal of Non-Linear Mechanics</i> , 2022, 140, 103907.	1.4	3
74	Models of Low Reynolds Number Swimmers Inspired by Cell Blebbing. <i>The IMA Volumes in Mathematics and Its Applications</i> , 2012, , 185-195.	0.5	2
75	The Status of the QSSA Approximation in Stochastic Simulations of Reaction Networks. <i>MATRIX Book Series</i> , 2020, , 137-147.	0.2	2
76	A Model for Pattern Formation in <i>Dictyostelium Discoideum</i> . <i>Lecture Notes in Biomathematics</i> , 1987, , 224-233.	0.3	2
77	Actin Cytoskeleton, Multi-scale Modeling. , 2015, , 17-23.		1
78	Cell-Based, Continuum and Hybrid Models of Tissue Dynamics. <i>Lecture Notes in Mathematics</i> , 2016, , 1-72.	0.1	1
79	A Random Walk Approach to Transport in Tissues and Complex Media: From Microscale Descriptions to Macroscale Models. <i>Bulletin of Mathematical Biology</i> , 2021, 83, 92.	0.9	1
80	Leveraging Compute Clusters for Large-Scale Parametric Screens of Reaction-Diffusion Systems. , 2018, , .		0