

Anotida Madzvamuse

List of Publications by Year in descending order

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Version: 2024-02-01

70
papers

1,710
citations

331670

21
h-index

315739

38
g-index

78
all docs

78
docs citations

78
times ranked

1212
citing authors

#	ARTICLE	IF	CITATIONS
1	Time-stepping schemes for moving grid finite elements applied to reaction–diffusion systems on fixed and growing domains. <i>Journal of Computational Physics</i> , 2006, 214, 239-263.	3.8	98
2	A moving grid finite element method applied to a model biological pattern generator. <i>Journal of Computational Physics</i> , 2003, 190, 478-500.	3.8	96
3	Stability analysis of non-autonomous reaction-diffusion systems: the effects of growing domains. <i>Journal of Mathematical Biology</i> , 2010, 61, 133-164.	1.9	89
4	A sensor kinase controls turgor-driven plant infection by the rice blast fungus. <i>Nature</i> , 2019, 574, 423-427.	27.8	87
5	The surface finite element method for pattern formation on evolving biological surfaces. <i>Journal of Mathematical Biology</i> , 2011, 63, 1095-1119.	1.9	84
6	Cross-diffusion-driven instability for reaction-diffusion systems: analysis and simulations. <i>Journal of Mathematical Biology</i> , 2015, 70, 709-743.	1.9	76
7	Velocity-induced numerical solutions of reaction-diffusion systems on continuously growing domains. <i>Journal of Computational Physics</i> , 2007, 225, 100-119.	3.8	72
8	A Moving Grid Finite Element Method for the Simulation of Pattern Generation by Turing Models on Growing Domains. <i>Journal of Scientific Computing</i> , 2005, 24, 247-262.	2.3	65
9	Two Interlinked Bistable Switches Govern Mitotic Control in Mammalian Cells. <i>Current Biology</i> , 2018, 28, 3824-3832.e6.	3.9	62
10	Cell migration through three-dimensional confining pores: speed accelerations by deformation and recoil of the nucleus. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2019, 374, 20180225.	4.0	62
11	A model for colour pattern formation in the butterfly wing of <i>Papilio dardanus</i> . <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2000, 267, 851-859.	2.6	57
12	Pigmentation pattern formation in butterflies: experiments and models. <i>Comptes Rendus - Biologies</i> , 2003, 326, 717-727.	0.2	56
13	A coupled bulk-surface model for cell polarisation. <i>Journal of Theoretical Biology</i> , 2019, 481, 119-135.	1.7	48
14	A Numerical Approach to the Study of Spatial Pattern Formation in the Ligaments of Arcoid Bivalves. <i>Bulletin of Mathematical Biology</i> , 2002, 64, 501-530.	1.9	41
15	A mechanochemical model of striae distensae. <i>Mathematical Biosciences</i> , 2012, 240, 141-147.	1.9	40
16	Stability analysis and simulations of coupled bulk-surface reaction–diffusion systems. <i>Proceedings of the Royal Society A: Mathematical, Physical and Engineering Sciences</i> , 2015, 471, 20140546.	2.1	39
17	Optogenetic Tuning Reveals Rho Amplification-Dependent Dynamics of a Cell Contraction Signal Network. <i>Cell Reports</i> , 2020, 33, 108467.	6.4	31
18	Implicit–Explicit Timestepping with Finite Element Approximation of Reaction–Diffusion Systems on Evolving Domains. <i>SIAM Journal on Numerical Analysis</i> , 2013, 51, 2309-2330.	2.3	29

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19	A Hybrid Multiscale Model for Cancer Invasion of the Extracellular Matrix. <i>Multiscale Modeling and Simulation</i> , 2020, 18, 824-850.	1.6	26
20	The bulk-surface finite element method for reaction-diffusion systems on stationary volumes. <i>Finite Elements in Analysis and Design</i> , 2016, 108, 9-21.	3.2	24
21	High Accuracy Benchmark Problems for Allen-Cahn and Cahn-Hilliard Dynamics. <i>Communications in Computational Physics</i> , 2019, 26, 947-972.	1.7	24
22	Global existence for semilinear reaction-diffusion systems on evolving domains. <i>Journal of Mathematical Biology</i> , 2012, 64, 41-67.	1.9	23
23	Dynamics of Shadow System of a Singular Gierer-Meinhardt System on an Evolving Domain. <i>Journal of Nonlinear Science</i> , 2021, 31, 1.	2.1	23
24	Modeling pattern formation during the early development of Amago trout. <i>Physical Review E</i> , 2011, 84, 041923.	2.1	22
25	Mathematical modelling and numerical simulations of actin dynamics in the eukaryotic cell. <i>Journal of Mathematical Biology</i> , 2013, 66, 547-593.	1.9	22
26	Exhibiting cross-diffusion-induced patterns for reaction-diffusion systems on evolving domains and surfaces. <i>Physical Review E</i> , 2014, 90, 043307.	2.1	20
27	Predicting and forecasting the impact of local outbreaks of COVID-19: use of SEIR-D quantitative epidemiological modelling for healthcare demand and capacity. <i>International Journal of Epidemiology</i> , 2021, 50, 1103-1113.	1.9	20
28	The moving grid finite element method applied to cell movement and deformation. <i>Finite Elements in Analysis and Design</i> , 2013, 74, 76-92.	3.2	19
29	Characterization of Turing diffusion-driven instability on evolving domains. <i>Discrete and Continuous Dynamical Systems</i> , 2012, 32, 3975-4000.	0.9	18
30	Stability analysis of reaction-diffusion models on evolving domains: The effects of cross-diffusion. <i>Discrete and Continuous Dynamical Systems</i> , 2015, 36, 2133-2170.	0.9	18
31	Stability analysis of Reaction-Diffusion Systems with constant coefficients on growing domains. <i>International Journal of Dynamical Systems and Differential Equations</i> , 2008, 1, 250.	0.0	17
32	Keratin Dynamics: Modeling the Interplay between Turnover and Transport. <i>PLoS ONE</i> , 2015, 10, e0121090.	2.5	16
33	Fully implicit time-stepping schemes and non-linear solvers for systems of reaction-diffusion equations. <i>Applied Mathematics and Computation</i> , 2014, 244, 361-374.	2.2	15
34	Lumped finite elements for reaction-cross-diffusion systems on stationary surfaces. <i>Computers and Mathematics With Applications</i> , 2017, 74, 3008-3023.	2.7	15
35	Bayesian Parameter Identification for Turing Systems on Stationary and Evolving Domains. <i>Bulletin of Mathematical Biology</i> , 2019, 81, 81-104.	1.9	15
36	Turing instability conditions for growing domains with divergence free mesh velocity. <i>Nonlinear Analysis: Theory, Methods & Applications</i> , 2009, 71, e2250-e2257.	1.1	12

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37	Analysis of stability and convergence of finite-difference methods for a reaction-diffusion problem on a one-dimensional growing domain. <i>IMA Journal of Numerical Analysis</i> , 2011, 31, 212-232.	2.9	12
38	A Model for Selection of Eyespots on Butterfly Wings. <i>PLoS ONE</i> , 2015, 10, e0141434.	2.5	12
39	A predictive model for color pattern formation in the butterfly wing of {it Papilio dardanus}. <i>Hiroshima Mathematical Journal</i> , 2002, 32, 325.	0.3	11
40	Classification of parameter spaces for a reaction-diffusion model on stationary domains. <i>Chaos, Solitons and Fractals</i> , 2017, 103, 33-51.	5.1	11
41	Bulk-surface virtual element method for systems of PDEs in two-space dimensions. <i>Numerische Mathematik</i> , 2021, 147, 305-348.	1.9	11
42	Wave of chaos in a spatial eco-epidemiological system: Generating realistic patterns of patchiness in rabbitâ€“lynx dynamics. <i>Mathematical Biosciences</i> , 2016, 281, 98-119.	1.9	10
43	Preserving invariance properties of reactionâ€“diffusion systems on stationary surfaces. <i>IMA Journal of Numerical Analysis</i> , 2019, 39, 235-270.	2.9	10
44	Whole cell tracking through the optimal control of geometric evolution laws. <i>Journal of Computational Physics</i> , 2015, 297, 495-514.	3.8	9
45	Projected finite elements for reactionâ€“diffusion systems on stationary closed surfaces. <i>Applied Numerical Mathematics</i> , 2015, 96, 45-71.	2.1	9
46	Numerical Preservation of Velocity Induced Invariant Regions for Reactionâ€“Diffusion Systems on Evolving Surfaces. <i>Journal of Scientific Computing</i> , 2018, 77, 971-1000.	2.3	9
47	An integrated framework for quantifying immune-tumour interactions in a 3D co-culture model. <i>Communications Biology</i> , 2021, 4, 781.	4.4	9
48	Growth patterns of noetiid ligaments: implications of developmental models for the origin of an evolutionary novelty among arcoid bivalves. <i>Geological Society Special Publication</i> , 2000, 177, 279-289.	1.3	8
49	A computational framework for particle and whole cell tracking applied to a real biological dataset. <i>Journal of Biomechanics</i> , 2016, 49, 1290-1304.	2.1	8
50	A Robust and Efficient Adaptive Multigrid Solver for the Optimal Control of Phase Field Formulations of Geometric Evolution Laws. <i>Communications in Computational Physics</i> , 2017, 21, 65-92.	1.7	8
51	The Stability Analyses of the Mathematical Models of Hepatitis C Virus Infection. <i>Modern Applied Science</i> , 2014, 9, .	0.6	7
52	A moving grid finite element method applied to a mechanobiochemical model for 3D cell migration. <i>Applied Numerical Mathematics</i> , 2020, 158, 336-359.	2.1	7
53	Projected Finite Elements for Systems of Reaction-Diffusion Equations on Closed Evolving Spheroidal Surfaces. <i>Communications in Computational Physics</i> , 2017, 21, 718-747.	1.7	6
54	Parameter identification through mode isolation for reactionâ€“diffusion systems on arbitrary geometries. <i>International Journal of Biomathematics</i> , 2018, 11, 1850053.	2.9	6

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55	From the Cell Membrane to the Nucleus: Unearthing Transport Mechanisms for Dynein. <i>Bulletin of Mathematical Biology</i> , 2012, 74, 2032-2061.	1.9	5
56	A mathematical understanding of how cytoplasmic dynein walks on microtubules. <i>Royal Society Open Science</i> , 2018, 5, 171568.	2.4	5
57	Domain-Dependent Stability Analysis of a Reaction–Diffusion Model on Compact Circular Geometries. <i>International Journal of Bifurcation and Chaos in Applied Sciences and Engineering</i> , 2018, 28, 1830024.	1.7	4
58	A mathematical analysis of an activator-inhibitor Rho GTPase model. <i>Journal of Computational Dynamics</i> , 2022, 9, 133.	1.1	4
59	Analysis and Simulations of Coupled Bulk-surface Reaction-Diffusion Systems on Exponentially Evolving Volumes. <i>Mathematical Modelling of Natural Phenomena</i> , 2016, 11, 4-32.	2.4	3
60	The Moving Grid Finite Element Method Applied to Biological Problems. , 2003, , 59-65.		3
61	Investigating Optimal Time Step Intervals of Imaging for Data Quality through a Novel Fully-Automated Cell Tracking Approach. <i>Journal of Imaging</i> , 2020, 6, 66.	3.0	2
62	Integrating Actin and Myosin II in a Viscous Model for Cell Migration. <i>Frontiers in Applied Mathematics and Statistics</i> , 2020, 6, .	1.3	2
63	Turing Pattern Formation Under Heterogeneous Distributions of Parameters for an Activator-Depleted Reaction Model. <i>Journal of Nonlinear Science</i> , 2021, 31, 1.	2.1	2
64	Domain-growth-induced patterning for reaction-diffusion systems with linear cross-diffusion. <i>Discrete and Continuous Dynamical Systems - Series B</i> , 2018, 23, 2775-2801.	0.9	2
65	Stability Analysis and Parameter Classification of a Reaction-Diffusion Model on an Annulus. <i>Journal of Applied Nonlinear Dynamics</i> , 2020, 9, 589-617.	0.3	2
66	Characterizing the Effects of Self- and Cross-Diffusion on Stationary Patterns of a Predator–Prey System. <i>International Journal of Bifurcation and Chaos in Applied Sciences and Engineering</i> , 2020, 30, 2050041.	1.7	1
67	Force Estimation during Cell Migration Using Mathematical Modelling. <i>Journal of Imaging</i> , 2022, 8, 199.	3.0	1
68	A Model for the Proliferation–Quiescence Transition in Human Cells. <i>Mathematics</i> , 2022, 10, 2426.	2.2	1
69	A note on how to develop interdisciplinary collaborations between experimentalists and theoreticians. <i>Interface Focus</i> , 2016, 6, 20160069.	3.0	0
70	Cross-Diffusion in Reaction-Diffusion Models: Analysis, Numerics, and Applications. <i>Mathematics in Industry</i> , 2017, , 385-392.	0.3	0