

Tomasz Lipniacki

List of Publications by Year in descending order

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Version: 2024-02-01

80
papers

3,147
citations

201575

27
h-index

175177

52
g-index

93
all docs

93
docs citations

93
times ranked

4537
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | The Spread of SARS-CoV-2 Variant Omicron with a Doubling Time of 2.0–3.3 Days Can Be Explained by Immune Evasion. <i>Viruses</i> , 2022, 14, 294. | 1.5 | 85 |
| 2 | Slow nucleosome dynamics set the transcriptional speed limit and induce RNA polymerase II traffic jams and bursts. <i>PLoS Computational Biology</i> , 2022, 18, e1009811. | 1.5 | 7 |
| 3 | Pareto-based evaluation of national responses to COVID-19 pandemic shows that saving lives and protecting economy are non-trade-off objectives. <i>Scientific Reports</i> , 2021, 11, 2425. | 1.6 | 28 |
| 4 | SARS-CoV-2 Variant of Concern 202012/01 Has about Twofold Replicative Advantage and Acquires Concerning Mutations. <i>Viruses</i> , 2021, 13, 392. | 1.5 | 92 |
| 5 | Traveling and standing fronts on curved surfaces. <i>Physica D: Nonlinear Phenomena</i> , 2020, 401, 132215. | 1.3 | 2 |
| 6 | A shear stress micromodel of urinary tract infection by the <i>Escherichia coli</i> producing Dr adhesin. <i>PLoS Pathogens</i> , 2020, 16, e1008247. | 2.1 | 16 |
| 7 | Electrochemical Immunosensors Based on Screen-Printed Gold and Glassy Carbon Electrodes: Comparison of Performance for Respiratory Syncytial Virus Detection. <i>Biosensors</i> , 2020, 10, 175. | 2.3 | 16 |
| 8 | Model-based optimization of combination protocols for irradiation-insensitive cancers. <i>Scientific Reports</i> , 2020, 10, 12652. | 1.6 | 2 |
| 9 | Super-spreading events initiated the exponential growth phase of COVID-19 with a higher than initially estimated. <i>Royal Society Open Science</i> , 2020, 7, 200786. | 1.1 | 47 |
| 10 | Coronavirus – Scientific insights and societal aspects. <i>Mathematical Modelling of Natural Phenomena</i> , 2020, 15, E2. | 0.9 | 15 |
| 11 | Dynamics of COVID-19 pandemic at constant and time-dependent contact rates. <i>Mathematical Modelling of Natural Phenomena</i> , 2020, 15, 28. | 0.9 | 41 |
| 12 | Modeling and measurement of signaling outcomes affecting decision making in noisy intracellular networks using machine learning methods. <i>Integrative Biology (United Kingdom)</i> , 2020, 12, 122-138. | 0.6 | 6 |
| 13 | A shear stress micromodel of urinary tract infection by the <i>Escherichia coli</i> producing Dr adhesin. , 2020, 16, e1008247. | | 0 |
| 14 | A shear stress micromodel of urinary tract infection by the <i>Escherichia coli</i> producing Dr adhesin. , 2020, 16, e1008247. | | 0 |
| 15 | A shear stress micromodel of urinary tract infection by the <i>Escherichia coli</i> producing Dr adhesin. , 2020, 16, e1008247. | | 0 |
| 16 | Robin-type boundary conditions in transition from reaction-diffusion equations in 3D domains to equations in 2D domains. <i>Journal of Differential Equations</i> , 2019, 268, 239-271. | 1.1 | 6 |
| 17 | Limits to the rate of information transmission through the MAPK pathway. <i>Journal of the Royal Society Interface</i> , 2019, 16, 20180792. | 1.5 | 10 |
| 18 | Cell fate in antiviral response arises in the crosstalk of IRF, NF- κ B and JAK/STAT pathways. <i>Nature Communications</i> , 2018, 9, 493. | 5.8 | 81 |

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|----|---|-----|-----------|
| 19 | Sampling rare events in stochastic reaction-diffusion systems within trajectory looping. <i>Physical Review E</i> , 2018, 98, 022401. | 0.8 | 1 |
| 20 | Relaxation oscillations and hierarchy of feedbacks in MAPK signaling. <i>Scientific Reports</i> , 2017, 7, 38244. | 1.6 | 47 |
| 21 | RAF1/BRAF dimerization integrates the signal from RAS to ERK and ROK1±. <i>Science Signaling</i> , 2017, 10, . | 1.6 | 40 |
| 22 | SPATKIN: a simulator for rule-based modeling of biomolecular site dynamics on surfaces. <i>Bioinformatics</i> , 2017, 33, 3667-3669. | 1.8 | 3 |
| 23 | Information processing in the NF- κ B pathway. <i>Scientific Reports</i> , 2017, 7, 15926. | 1.6 | 25 |
| 24 | Polarization of concave domains by traveling wave pinning. <i>PLoS ONE</i> , 2017, 12, e0190372. | 1.1 | 4 |
| 25 | Genetic toggle switch controlled by bacterial growth rate. <i>BMC Systems Biology</i> , 2017, 11, 117. | 3.0 | 18 |
| 26 | Computation and measurement of cell decision making errors using single cell data. <i>PLoS Computational Biology</i> , 2017, 13, e1005436. | 1.5 | 18 |
| 27 | Importins promote high-frequency NF- κ B oscillations increasing information channel capacity. <i>Biology Direct</i> , 2016, 11, 61. | 1.9 | 15 |
| 28 | Feedbacks, Bifurcations, and Cell Fate Decision-Making in the p53 System. <i>PLoS Computational Biology</i> , 2016, 12, e1004787. | 1.5 | 46 |
| 29 | Clustering reveals limits of parameter identifiability in multi-parameter models of biochemical dynamics. <i>BMC Systems Biology</i> , 2015, 9, 65. | 3.0 | 17 |
| 30 | Effective reaction rates for diffusion-limited reaction cycles. <i>Journal of Chemical Physics</i> , 2015, 143, 215102. | 1.2 | 4 |
| 31 | Computational Analysis of an Autophagy/Translation Switch Based on Mutual Inhibition of MTORC1 and ULK1. <i>PLoS ONE</i> , 2015, 10, e0116550. | 1.1 | 38 |
| 32 | Effective reaction rates in diffusion-limited phosphorylation-dephosphorylation cycles. <i>Physical Review E</i> , 2015, 91, 022702. | 0.8 | 7 |
| 33 | Digital signaling decouples activation probability and population heterogeneity. <i>ELife</i> , 2015, 4, e08931. | 2.8 | 60 |
| 34 | Stability of bacterial toggle switches is enhanced by cell-cycle lengthening by several orders of magnitude. <i>Physical Review E</i> , 2014, 89, 022710. | 0.8 | 8 |
| 35 | Dynamic Cross Talk Model of the Epithelial Innate Immune Response to Double-Stranded RNA Stimulation: Coordinated Dynamics Emerging from Cell-Level Noise. <i>PLoS ONE</i> , 2014, 9, e93396. | 1.1 | 33 |
| 36 | A Spatially Extended Model of Kinase-Receptor Interaction. <i>SIAM Journal on Applied Mathematics</i> , 2013, 73, 374-400. | 0.8 | 5 |

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|----|--|------|-----------|
| 37 | Levels of pro-apoptotic regulator Bad and anti-apoptotic regulator Bcl-xL determine the type of the apoptotic logic gate. <i>BMC Systems Biology</i> , 2013, 7, 67. | 3.0 | 29 |
| 38 | Type of noise defines global attractors in bistable molecular regulatory systems. <i>Journal of Theoretical Biology</i> , 2013, 317, 140-151. | 0.8 | 23 |
| 39 | Stochastic transitions in a bistable reaction system on the membrane. <i>Journal of the Royal Society Interface</i> , 2013, 10, 20130151. | 1.5 | 14 |
| 40 | MEK1 and MEK2 differentially control the duration and amplitude of the ERK cascade response. <i>Physical Biology</i> , 2013, 10, 035006. | 0.8 | 11 |
| 41 | Toggle switch: noise determines the winning gene. <i>Physical Biology</i> , 2013, 10, 035007. | 0.8 | 20 |
| 42 | Spontaneous NF- κ B Activation by Autocrine TNF α Signaling: A Computational Analysis. <i>PLoS ONE</i> , 2013, 8, e78887. | 1.1 | 57 |
| 43 | A Computational Model for Early Events in B Cell Antigen Receptor Signaling: Analysis of the Roles of Lyn and Fyn. <i>Journal of Immunology</i> , 2012, 189, 646-658. | 0.4 | 46 |
| 44 | Dynamics of a stochastic spatially extended system predicted by comparing deterministic and stochastic attractors of the corresponding birth-death process. <i>Physical Biology</i> , 2012, 9, 055002. | 0.8 | 13 |
| 45 | The interplay of double phosphorylation and scaffolding in MAPK pathways. <i>Journal of Theoretical Biology</i> , 2012, 295, 116-124. | 0.8 | 31 |
| 46 | Guidelines for visualizing and annotating rule-based models. <i>Molecular BioSystems</i> , 2011, 7, 2779. | 2.9 | 36 |
| 47 | Vortex loops cascade as a channel of quantum turbulence decay. <i>Journal of Physics: Conference Series</i> , 2011, 318, 092028. | 0.3 | 0 |
| 48 | Cascade of vortex loops initiated by a single reconnection of quantum vortices. <i>Physical Review B</i> , 2011, 83, . | 1.1 | 45 |
| 49 | Exact solutions to a spatially extended model of kinase-receptor interaction. <i>Physical Biology</i> , 2011, 8, 055005. | 0.8 | 3 |
| 50 | B Cell Activation Triggered by the Formation of the Small Receptor Cluster: A Computational Study. <i>PLoS Computational Biology</i> , 2011, 7, e1002197. | 1.5 | 29 |
| 51 | Spatial gradients in kinase cascade regulation. <i>IET Systems Biology</i> , 2010, 4, 348-355. | 0.8 | 13 |
| 52 | Single-cell NF- κ B dynamics reveal digital activation and analogue information processing. <i>Nature</i> , 2010, 466, 267-271. | 13.7 | 736 |
| 53 | Exploring mechanisms of oscillations in p53 and nuclear factor- κ B systems. <i>IET Systems Biology</i> , 2009, 3, 342-355. | 0.8 | 25 |
| 54 | Crosstalk between p53 and nuclear factor- κ B systems: pro- and anti-apoptotic functions of NF- κ B. <i>IET Systems Biology</i> , 2009, 3, 356-367. | 0.8 | 45 |

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|----|--|-----|-----------|
| 55 | Regulation of kinase activity by diffusion and feedback. <i>Journal of Theoretical Biology</i> , 2009, 259, 291-296. | 0.8 | 33 |
| 56 | Stochastic effects and bistability in T cell receptor signaling. <i>Journal of Theoretical Biology</i> , 2008, 254, 110-122. | 0.8 | 86 |
| 57 | Oscillations and bistability in the stochastic model of p53 regulation. <i>Journal of Theoretical Biology</i> , 2008, 254, 452-465. | 0.8 | 108 |
| 58 | Deterministic and Stochastic Models of NF κ B Pathway. <i>Cardiovascular Toxicology</i> , 2007, 7, 215-234. | 1.1 | 23 |
| 59 | Adjoint Systems for Models of Cell Signaling Pathways and their Application to Parameter Fitting. <i>IEEE/ACM Transactions on Computational Biology and Bioinformatics</i> , 2007, 4, 322-335. | 1.9 | 26 |
| 60 | Single TNF α trimers mediating NF- κ B activation: stochastic robustness of NF- κ B signaling. <i>BMC Bioinformatics</i> , 2007, 8, 376. | 1.2 | 60 |
| 61 | How the Number of Alleles Influences Gene Expression. <i>Journal of Statistical Physics</i> , 2007, 128, 511-533. | 0.5 | 13 |
| 62 | Asymptotic behavior of distributions of mRNA and protein levels in a model of stochastic gene expression. <i>Journal of Mathematical Analysis and Applications</i> , 2007, 333, 753-769. | 0.5 | 45 |
| 63 | Stochastic Regulation in Early Immune Response. <i>Biophysical Journal</i> , 2006, 90, 725-742. | 0.2 | 86 |
| 64 | Dynamics of superfluid 4He: Two-scale approach. <i>European Journal of Mechanics, B/Fluids</i> , 2006, 25, 435-458. | 1.2 | 15 |
| 65 | Transcriptional stochasticity in gene expression. <i>Journal of Theoretical Biology</i> , 2006, 238, 348-367. | 0.8 | 120 |
| 66 | Stochastic effects of multiple regulators on expression profiles in eukaryotes. <i>Journal of Theoretical Biology</i> , 2005, 233, 423-433. | 0.8 | 24 |
| 67 | Mathematical model of NF- κ B regulatory module. <i>Journal of Theoretical Biology</i> , 2004, 228, 195-215. | 0.8 | 264 |
| 68 | Quasi-static solutions for quantum vortex motion under the localized induction approximation. <i>Journal of Fluid Mechanics</i> , 2003, 477, . | 1.4 | 25 |
| 69 | Shape-preserving solutions for quantum vortex motion under localized induction approximation. <i>Physics of Fluids</i> , 2003, 15, 1381. | 1.6 | 23 |
| 70 | Homoclinic solutions in mechanical systems with small dissipation. Application to DNA dynamics. <i>Journal of Mathematical Biology</i> , 2002, 44, 309-329. | 0.8 | 4 |
| 71 | Evolution of the anisotropy of the quantum vortex tangle. , 2002, , 93-98. | | 0 |
| 72 | Torsional Travelling Waves in DNA. <i>Journal of Nonlinear Mathematical Physics</i> , 2001, 8, 188. | 0.8 | 0 |

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|----|--|-----|-----------|
| 73 | STATICS OF RIGID UNITS CHAIN. International Journal of Bifurcation and Chaos in Applied Sciences and Engineering, 2001, 11, 845-855. | 0.7 | 0 |
| 74 | Thermodynamics of local DNA openings. Physical Review E, 2001, 64, 051919. | 0.8 | 15 |
| 75 | Evolution of the line-length density and anisotropy of quantum tangle in ^4He . Physical Review B, 2001, 64, . | 1.1 | 20 |
| 76 | From Vortex Reconnections to Quantum Turbulence. , 2001, , 177-183. | | 2 |
| 77 | Torsional Travelling Waves in DNA. Journal of Nonlinear Mathematical Physics, 2001, 8, 188. | 0.8 | 0 |
| 78 | Evolution of quantum vortices following reconnection. European Journal of Mechanics, B/Fluids, 2000, 19, 361-378. | 1.2 | 22 |
| 79 | Chemically driven traveling waves in DNA. Physical Review E, 1999, 60, 7253-7261. | 0.8 | 21 |
| 80 | Non-linear mechanical model of DNA dynamics. Nuovo Cimento Della Societa Italiana Di Fisica D - Condensed Matter, Atomic, Molecular and Chemical Physics, Biophysics, 1998, 20, 833-843. | 0.4 | 7 |