

Katarzyna A Rejniak

List of Publications by Year in descending order

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Version: 2024-02-01

43
papers

1,900
citations

304602

22
h-index

276775

41
g-index

50
all docs

50
docs citations

50
times ranked

2114
citing authors

#	ARTICLE	IF	CITATIONS
1	Collective Cell Migration in a Fibrous Environment: A Hybrid Multiscale Modelling Approach. <i>Frontiers in Applied Mathematics and Statistics</i> , 2021, 7, .	0.7	15
2	Bridging cell-scale simulations and radiologic images to explain short-time intratumoral oxygen fluctuations. <i>PLoS Computational Biology</i> , 2021, 17, e1009206.	1.5	7
3	Hybrid modeling frameworks of tumor development and treatment. <i>Wiley Interdisciplinary Reviews: Systems Biology and Medicine</i> , 2020, 12, e1461.	6.6	56
4	High School Internship Program in Integrated Mathematical Oncology (HIP IMO): Five-Year Experience at Moffitt Cancer Center. <i>Bulletin of Mathematical Biology</i> , 2020, 82, 91.	0.9	4
5	Comparison of Drug Inhibitory Effects (IC_{50}) in Monolayer and Spheroid Cultures. <i>Bulletin of Mathematical Biology</i> , 2020, 82, 68.	0.9	46
6	Drug-Induced Resistance in Micrometastases: Analysis of Spatio-Temporal Cell Lineages. <i>Frontiers in Physiology</i> , 2020, 11, 319.	1.3	21
7	Morphophenotypic classification of tumor organoids as an indicator of drug exposure and penetration potential. <i>PLoS Computational Biology</i> , 2019, 15, e1007214.	1.5	24
8	Single-Cell-Based In Silico Models: A Tool for Dissecting Tumor Heterogeneity. , 2019, , 130-143.		6
9	Micropharmacology: An In Silico Approach for Assessing Drug Efficacy Within a Tumor Tissue. <i>Bulletin of Mathematical Biology</i> , 2019, 81, 3623-3641.	0.9	19
10	Towards personalized computational oncology: from spatial models of tumour spheroids, to organoids, to tissues. <i>Journal of the Royal Society Interface</i> , 2018, 15, 20170703.	1.5	101
11	Targeting Ligand Specificity Linked to Tumor Tissue Topological Heterogeneity via Single-Cell Micro-Pharmacological Modeling. <i>Scientific Reports</i> , 2018, 8, 3638.	1.6	12
12	Mathematical Modeling of Tumor Organoids: Toward Personalized Medicine. <i>Cancer Drug Discovery and Development</i> , 2018, , 193-213.	0.2	1
13	Circulating Tumor Cells: When a Solid Tumor Meets a Fluid Microenvironment. <i>Advances in Experimental Medicine and Biology</i> , 2016, 936, 93-106.	0.8	68
14	Microenvironmental Niches and Sanctuaries: A Route to Acquired Resistance. <i>Advances in Experimental Medicine and Biology</i> , 2016, 936, 149-164.	0.8	24
15	Limiting the development of anti-cancer drug resistance in a spatial model of micrometastases. <i>Mathematical Biosciences and Engineering</i> , 2016, 13, 1185-1206.	1.0	26
16	Pathology to Enhance Precision Medicine in Oncology. <i>Advances in Anatomic Pathology</i> , 2015, 22, 267-272.	2.4	30
17	Diagnostic assessment of osteosarcoma chemoresistance based on Virtual Clinical Trials. <i>Medical Hypotheses</i> , 2015, 85, 348-354.	0.8	24
18	Pyruvate sensitizes pancreatic tumors to hypoxia-activated prodrug TH-302. <i>Cancer & Metabolism</i> , 2015, 3, 2.	2.4	69

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19	Emergence of Anti-Cancer Drug Resistance: Exploring the Importance of the Microenvironmental Niche via a Spatial Model. <i>The IMA Volumes in Mathematics and Its Applications</i> , 2015, , 1-34.	0.5	16
20	Mechanical Aspects of Microtubule Bundling in Taxane-Treated Circulating Tumor Cells. <i>Biophysical Journal</i> , 2014, 107, 1236-1246.	0.2	9
21	The formation of tight tumor clusters affects the efficacy of cell cycle inhibitors: A hybrid model study. <i>Journal of Theoretical Biology</i> , 2014, 352, 31-50.	0.8	26
22	On a conditionally stable nonlinear method to approximate some monotone and bounded solutions of a generalized population model. <i>Applied Mathematics and Computation</i> , 2014, 229, 273-282.	1.4	5
23	IBCell Morphocharts: A Computational Model for Linking Cell Molecular Activity with Emerging Tissue Morphology. <i>Natural Computing Series</i> , 2014, , 507-524.	2.2	3
24	Current Advances in Mathematical Modeling of Anti-Cancer Drug Penetration into Tumor Tissues. <i>Frontiers in Oncology</i> , 2013, 3, 278.	1.3	102
25	The Role of Tumor Tissue Architecture in Treatment Penetration and Efficacy: An Integrative Study. <i>Frontiers in Oncology</i> , 2013, 3, 111.	1.3	62
26	Simulating Cancer: Computational Models in Oncology. <i>Frontiers in Oncology</i> , 2013, 3, 233.	1.3	23
27	Computational investigation of intrinsic and extrinsic mechanisms underlying the formation of carcinoma. <i>Mathematical Medicine and Biology</i> , 2012, 29, 67-84.	0.8	15
28	Homeostatic Imbalance in Epithelial Ducts and Its Role in Carcinogenesis. <i>Scientifica</i> , 2012, 2012, 1-8.	0.6	9
29	Investigating dynamical deformations of tumor cells in circulation: predictions from a theoretical model. <i>Frontiers in Oncology</i> , 2012, 2, 111.	1.3	40
30	Cellular modeling of cancer invasion: Integration of in silico and in vitro approaches. <i>Journal of Cellular Physiology</i> , 2012, 227, 431-438.	2.0	68
31	Hybrid models of tumor growth. <i>Wiley Interdisciplinary Reviews: Systems Biology and Medicine</i> , 2011, 3, 115-125.	6.6	256
32	Current trends in mathematical modeling of tumorâ€™microenvironment interactions: a survey of tools and applications. <i>Experimental Biology and Medicine</i> , 2010, 235, 411-423.	1.1	54
33	Linking Changes in Epithelial Morphogenesis to Cancer Mutations Using Computational Modeling. <i>PLoS Computational Biology</i> , 2010, 6, e1000900.	1.5	38
34	Microenvironment driven invasion: a multiscale multimodel investigation. <i>Journal of Mathematical Biology</i> , 2009, 58, 579-624.	0.8	92
35	A Computational Study of the Development of Epithelial Acini: I. Sufficient Conditions for the Formation of a Hollow Structure. <i>Bulletin of Mathematical Biology</i> , 2008, 70, 677-712.	0.9	54
36	A Computational Study of the Development of Epithelial Acini: II. Necessary Conditions for Structure and Lumen Stability. <i>Bulletin of Mathematical Biology</i> , 2008, 70, 1450-1479.	0.9	36

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37	Invasion emerges from cancer cell adaptation to competitive microenvironments: Quantitative predictions from multiscale mathematical models. <i>Seminars in Cancer Biology</i> , 2008, 18, 338-348.	4.3	64
38	A Single Cell-Based Model of the Ductal Tumour Microarchitecture. <i>Computational and Mathematical Methods in Medicine</i> , 2007, 8, 51-69.	0.7	63
39	Modelling the Development of Complex Tissues Using Individual Viscoelastic Cells. , 2007, , 301-323.		11
40	An immersed boundary framework for modelling the growth of individual cells: An application to the early tumour development. <i>Journal of Theoretical Biology</i> , 2007, 247, 186-204.	0.8	156
41	A Single-Cell Approach in Modeling the Dynamics of Tumor Microregions. <i>Mathematical Biosciences and Engineering</i> , 2005, 2, 643-655.	1.0	68
42	Digital Transcriptome Analysis in the Aging Cerebellum. <i>Annals of the New York Academy of Sciences</i> , 2004, 1019, 58-63.	1.8	7
43	A computational model of the mechanics of growth of the villous trophoblast bilayer. <i>Bulletin of Mathematical Biology</i> , 2004, 66, 199-232.	0.9	50