Igor Goryanin

List of Publications by Year in descending order

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71532 236612 7,093 81 25 76 citations h-index g-index papers 85 85 85 7944 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	The systems biology markup language (SBML): a medium for representation and exchange of biochemical network models. Bioinformatics, 2003, 19, 524-531.	1.8	2,811
2	A community-driven global reconstruction of human metabolism. Nature Biotechnology, 2013, 31, 419-425.	9.4	920
3	The Systems Biology Graphical Notation. Nature Biotechnology, 2009, 27, 735-741.	9.4	828
4	The Edinburgh human metabolic network reconstruction and its functional analysis. Molecular Systems Biology, 2007, 3, 135.	3.2	364
5	Metabolic modeling of microbial strains in silico. Trends in Biochemical Sciences, 2001, 26, 179-186.	3.7	291
6	Novel Electrochemically Active Bacterium Phylogenetically Related to <i>Arcobacter butzleri</i> , Isolated from a Microbial Fuel Cell. Applied and Environmental Microbiology, 2009, 75, 7326-7334.	1.4	169
7	Systems Biology Reveals New Strategies for Personalizing Cancer Medicine and Confirms the Role of PTEN in Resistance to Trastuzumab. Cancer Research, 2009, 69, 6713-6720.	0.4	152
8	Mathematical simulation and analysis of cellular metabolism and regulation. Bioinformatics, 1999, 15, 749-758.	1.8	135
9	A fragile metabolic network adapted for cooperation in the symbiotic bacterium Buchnera aphidicola. BMC Systems Biology, 2009, 3, 24.	3.0	98
10	Compartmentalization of the Edinburgh Human Metabolic Network. BMC Bioinformatics, 2010, 11, 393.	1.2	92
11	The metabolic pathway collection from EMP: the enzymes and metabolic pathways database. Nucleic Acids Research, 1996, 24, 26-28.	6.5	87
12	Temperature Dependence of the Epidermal Growth Factor Receptor Signaling Network Can Be Accounted for by a Kinetic Modelâ€. Biochemistry, 2002, 41, 306-320.	1.2	74
13	EchoBASE: an integrated post-genomic database for Escherichia coli. Nucleic Acids Research, 2004, 33, D329-D333.	6.5	70
14	Human metabolic network reconstruction and its impact on drug discovery and development. Drug Discovery Today, 2008, 13, 402-408.	3.2	52
15	Passive microwave radiometry in biomedical studies. Drug Discovery Today, 2020, 25, 757-763.	3.2	49
16	Mathematical Modeling of Mitochondrial Adenine Nucleotide Translocase. Biophysical Journal, 2006, 90, 423-432.	0.2	43
17	EnzML: multi-label prediction of enzyme classes using InterPro signatures. BMC Bioinformatics, 2012, 13, 61.	1.2	43
18	DBSolve Optimum: a software package for kinetic modeling which allows dynamic visualization of simulation results. BMC Systems Biology, 2010, 4, 109.	3.0	39

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19	The metabolic pathway collection: an update. Nucleic Acids Research, 1997, 25, 37-38.	6.5	37
20	Microarray data can predict diurnal changes of starch content in the picoalga Ostreococcus. BMC Systems Biology, 2011, 5, 36.	3.0	37
21	Kinetic Model of Mitochondrial Krebs Cycle: Unraveling the Mechanism of Salicylate Hepatotoxic Effects. Journal of Biological Physics, 2006, 32, 245-271.	0.7	36
22	Model-based global sensitivity analysis as applied to identification of anti-cancer drug targets and biomarkers of drug resistance in the ErbB2/3 network. European Journal of Pharmaceutical Sciences, 2012, 46, 244-258.	1.9	35
23	Towards a computational reconstruction of the electrodynamics of premature and full term human labour. Progress in Biophysics and Molecular Biology, 2011, 107, 183-192.	1.4	29
24	Pathway-Consensus Approach to Metabolic Network Reconstruction for Pseudomonas putida KT2440 by Systematic Comparison of Published Models. PLoS ONE, 2017, 12, e0169437.	1.1	29
25	Modern Microwave Thermometry for Breast Cancer. Journal of Molecular Imaging & Dynamics, 2017, 7,	0.2	27
26	Using medical microwave radiometry for brain temperature measurements. Drug Discovery Today, 2022, 27, 881-889.	3.2	26
27	The Pathway Editor: A tool for managing complex biological networks. IBM Journal of Research and Development, 2006, 50, 561-573.	3.2	25
28	The reconstruction and analysis of tissue specific human metabolic networks. Molecular BioSystems, 2012, 8, 663-670.	2.9	25
29	Kinetic Modeling of Energy Metabolism and Superoxide Generation in Hepatocyte Mitochondria. Molecular Biology, 2001, 35, 940-949.	0.4	23
30	Modelling nitrogen assimilation of Escherichia coli at low ammonium concentration. Journal of Biotechnology, 2009, 144, 175-183.	1.9	22
31	Multi-electrode microbial fuel cell with horizontal liquid flow. Water Science and Technology, 2009, 60, 347-355.	1.2	20
32	SBSI: an extensible distributed software infrastructure for parameter estimation in systems biology. Bioinformatics, 2013, 29, 664-665.	1.8	20
33	The kinetic model of the shikimate pathway as a tool to optimize enzyme assays for high-throughput screening. Biotechnology and Bioengineering, 2006, 95, 560-571.	1.7	19
34	Kinetic modelling of NSAID action on COX-1: Focus on in vitro/in vivo aspects and drug combinations. European Journal of Pharmaceutical Sciences, 2009, 36, 122-136.	1.9	19
35	Compensatory effects in the PI3K/PTEN/AKT signaling network following receptor tyrosine kinase inhibition. Cellular Signalling, 2011, 23, 407-416.	1.7	19
36	Single chamber air–cathode microbial fuel cells as biosensors for determination of biodegradable organics. Biotechnology Letters, 2019, 41, 555-563.	1.1	18

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37	The impact of the regulatory design on the response of epidermal growth factor receptorâ€mediated signal transduction towards oncogenic mutations. FEBS Journal, 2007, 274, 5505-5517.	2.2	15
38	A Graphical Notation to Describe the Logical Interactions of Biological Pathways. Journal of Integrative Bioinformatics, 2006, 3, 177-187.	1.0	14
39	Concurrent treatment of raw and aerated swine wastewater using an electrotrophic denitrification system. Bioresource Technology, 2021, 322, 124508.	4.8	14
40	Passive Microwave Radiometry for the Diagnosis of Coronavirus Disease 2019 Lung Complications in Kyrgyzstan. Diagnostics, 2021, 11, 259.	1.3	14
41	Simultaneous modelling of metabolic, genetic and product-interaction networks. Briefings in Bioinformatics, 2001, 2, 223-232.	3.2	13
42	Kinetic Modeling as a Tool to Integrate Multilevel Dynamic Experimental Data. Methods in Molecular Biology, 2009, 563, 197-218.	0.4	13
43	A semi-automated genome annotation comparison and integration scheme. BMC Bioinformatics, 2013, 14, 172.	1.2	13
44	Metabolic engineering of a novel strain of electrogenic bacterium Arcobacter butzleri to create a platform for single analyte detection using a microbial fuel cell. Enzyme and Microbial Technology, 2020, 139, 109564.	1.6	13
45	Application of Data Mining and Machine Learning in Microwave Radiometry (MWR). Communications in Computer and Information Science, 2020, , 265-288.	0.4	12
46	Application of Artificial Intelligence in Microwave Radiometry (MWR)., 2019,,.		12
47	KINETIC MODEL OF PHOSPHOFRUCTOKINASE-1 FROMESCHERICHIA COLI. Journal of Bioinformatics and Computational Biology, 2008, 06, 843-867.	0.3	11
48	Use of Microwave Radiometry to Monitor Thermal Denaturation of Albumin. Frontiers in Physiology, 2018, 9, 956.	1.3	11
49	Microwave Radiometry (MWR) temperature measurement is related to symptom severity in patients with Low Back Pain (LBP). Journal of Bodywork and Movement Therapies, 2021, 26, 548-552.	0.5	11
50	Using AI and passive medical radiometry for diagnostics (MWR) of venous diseases. Computer Methods and Programs in Biomedicine, 2022, 215, 106611.	2.6	11
51	Dynamic computational modeling in the search for better breast cancer drug therapy. Pharmacogenomics, 2007, 8, 1757-1761.	0.6	10
52	In Silico Screening of Nonsteroidal Anti-Inflammatory Drugs and Their Combined Action on Prostaglandin H Synthase-1. Pharmaceuticals, 2010, 3, 2059-2081.	1.7	9
53	Taxonomic and functional metagenomic analysis of anodic communities in two pilot-scale microbial fuel cells treating different industrial wastewaters. Journal of Integrative Bioinformatics, 2015, 12, 1-15.	1.0	9
54	Magnet-Facilitated Selection of Electrogenic Bacteria from Marine Sediment. BioMed Research International, 2015, 2015, 1-7.	0.9	9

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55	El of the Phosphotransferase System of Escherichia coli: Mathematical Modeling Approach to Analysis of Its Kinetic Properties. Journal of Biophysics, 2011, 2011, 1-17.	0.8	8
56	Comparative Metagenomic Analysis of Electrogenic Microbial Communities in Differentially Inoculated Swine Wastewater-Fed Microbial Fuel Cells. Scientifica, 2017, 2017, 1-10.	0.6	7
57	Computational optimization and biological evolution. Biochemical Society Transactions, 2010, 38, 1206-1209.	1.6	6
58	Modern microwave thermometry for breast cancer. , 2017, , .		6
59	Genome Sequence of the Electrogenic Petroleum-Degrading Thalassospira sp. Strain HJ. Genome Announcements, 2015, 3, .	0.8	5
60	ASAR: visual analysis of metagenomes in R. Bioinformatics, 2018, 34, 1404-1405.	1.8	5
61	Treatment and Companion Diagnostics of Lower Back Pain Using Self-Controlled Energo-Neuroadaptive Regulator (SCENAR) and Passive Microwave Radiometry (MWR). Diagnostics, 2022, 12, 1220.	1.3	5
62	A computer model of the gene network of the cholesterol biosynthesis regulation in the cell: analysis of the effect of mutations. Doklady Biochemistry and Biophysics, 2003, 389, 90-93.	0.3	4
63	Kinetic model of functioning and regulation of Escherichia coli isocitrate dehydrogenase. Biophysics (Russian Federation), 2007, 52, 30-39.	0.2	4
64	KINETIC MODELING OFACEOPERON GENETIC REGULATION INESCHERICHIA COLI. Journal of Bioinformatics and Computational Biology, 2008, 06, 933-959.	0.3	4
65	Taxonomic and functional metagenomic analysis of anodic communities in two pilot-scale microbial fuel cells treating different industrial wastewaters. Journal of Integrative Bioinformatics, 2015, 12, 273.	1.0	4
66	Passive Microwave Radiometry as a Component of Imaging Diagnostics in Juvenile Idiopathic Arthritis. Rheumato, 2022, 2, 55-68.	0.2	4
67	Reconstructing whole-cell models. Drug Discovery Today, 2001, 6, 109-112.	3.2	3
68	Kinetic model of imidazologlycerol-phosphate synthetase from Escherichia coli. Biochemistry (Moscow), 2004, 69, 1324-1335.	0.7	3
69	Preface. European Journal of Pharmaceutical Sciences, 2012, 46, 189.	1.9	3
70	Processivity and Coupling in Messenger RNA Transcription. PLoS ONE, 2010, 5, e8845.	1.1	3
71	A Method of Microwave Radiothermometry in Studies of Circadian Rhythms of Brain Temperature. Bulletin of Experimental Biology and Medicine, 2022, 173, 380-383.	0.3	3
72	Cellular Kinetic Modeling of the Microbial Metabolism. Methods of Biochemical Analysis, 2005, , 437-488.	0.2	2

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73	Is there a Function for a Sex Pheromone Precursor?. Journal of Integrative Bioinformatics, 2019, 16, .	1.0	2
74	Kinetic Modeling of E. coli Enzymes: Integration of in vitro Experimental Data., 2009, , 177-207.		2
75	Preface. European Journal of Pharmaceutical Sciences, 2009, 36, 1-3.	1.9	1
76	A kinetic model of Escherichia coli β-galactosidase. Biophysics (Russian Federation), 2009, 54, 156-162.	0.2	1
77	A user-defined data type for the storage of time series data allowing efficient similarity screening. European Journal of Pharmaceutical Sciences, 2012, 46, 272-274.	1.9	1
78	Bioinformatic analysis of bacterial composition and metabolic mapping of selectively enriched microbial community within Microbial Fuel Cells., 2017,,.		0
79	WikiSim., 2008,,.		0
80	ASAR Database: An R Tool for Visual Analysis and Storage of Metagenomes. , 2018, , .		0
81	Monitoring Protein Denaturation of Egg White Using Passive Microwave Radiometry (MWR). Diagnostics, 2022, 12, 1498.	1.3	O