

Chris John Myers

List of Publications by Year in descending order

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155
papers

3,306
citations

201385

27
h-index

243296

44
g-index

168
all docs

168
docs citations

168
times ranked

2339
citing authors

#	ARTICLE	IF	CITATIONS
1	Round Trip: An Automated Pipeline for Experimental Design, Execution, and Analysis. ACS Synthetic Biology, 2022, 11, 608-622.	1.9	8
2	Sequence-Based Searching for SynBioHub Using VSEARCH. ACS Synthetic Biology, 2022, 11, 990-995.	1.9	4
3	Genetic circuit design automation with Cello 2.0. Nature Protocols, 2022, 17, 1097-1113.	5.5	52
4	BioSimulators: a central registry of simulation engines and services for recommending specific tools. Nucleic Acids Research, 2022, 50, W108-W114.	6.5	11
5	Engineering genetic circuits: advancements in genetic design automation tools and standards for synthetic biology. Current Opinion in Microbiology, 2022, 68, 102155.	2.3	7
6	Synthetic biology open language visual (SBOL Visual) version 2.3. Journal of Integrative Bioinformatics, 2021, 18, .	1.0	6
7	SBOLCanvas: A Visual Editor for Genetic Designs. ACS Synthetic Biology, 2021, 10, 1792-1796.	1.9	9
8	Quantitative characterization of recombinase-based digitizer circuits enables predictable amplification of biological signals. Communications Biology, 2021, 4, 875.	2.0	9
9	VisBOL2â€”Improving Web-Based Visualization for Synthetic Biology Designs. ACS Synthetic Biology, 2021, 10, 2111-2115.	1.9	3
10	Synthetic Biology Knowledge System. ACS Synthetic Biology, 2021, 10, 2276-2285.	1.9	9
11	Curation Principles Derived from the Analysis of the SBOL iGEM Data Set. ACS Synthetic Biology, 2021, 10, 2592-2606.	1.9	3
12	Stochastic Hazard Analysis of Genetic Circuits in iBioSim and STAMINA. ACS Synthetic Biology, 2021, 10, 2532-2540.	1.9	2
13	Specifications of standards in systems and synthetic biology: status and developments in 2021. Journal of Integrative Bioinformatics, 2021, 18, .	1.0	2
14	Synthetic Biology Curation Tools (SYNBICT). ACS Synthetic Biology, 2021, 10, 3200-3204.	1.9	2
15	The Synthetic Biology Open Language (SBOL) Version 3: Simplified Data Exchange for Bioengineering. Frontiers in Bioengineering and Biotechnology, 2020, 8, 1009.	2.0	40
16	Genetic Circuit Dynamics: Hazard and Glitch Analysis. ACS Synthetic Biology, 2020, 9, 2324-2338.	1.9	21
17	SBOL Visual 2 Ontology. ACS Synthetic Biology, 2020, 9, 972-977.	1.9	3
18	Extending SynBioHubâ€™s Functionality with Plugins. ACS Synthetic Biology, 2020, 9, 1216-1220.	1.9	9

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19	The first 10 years of the international coordination network for standards in systems and synthetic biology (COMBINE). Journal of Integrative Bioinformatics, 2020, 17, .	1.0	18
20	Synthetic biology open language (SBOL) version 3.0.0. Journal of Integrative Bioinformatics, 2020, 17, .	1.0	13
21	Systems Biology Markup Language (SBML) Level 3 Package: Distributions, Version 1, Release 1. Journal of Integrative Bioinformatics, 2020, 17, .	1.0	7
22	Specifications of standards in systems and synthetic biology: status and developments in 2020. Journal of Integrative Bioinformatics, 2020, 17, .	1.0	10
23	The long journey towards standards for engineering biosystems. EMBO Reports, 2020, 21, e50521.	2.0	46
24	<sc>SBML</sc> Level 3: an extensible format for the exchange and reuse of biological models. Molecular Systems Biology, 2020, 16, e9110.	3.2	178
25	A Computational Workflow for the Automated Generation of Models of Genetic Designs. ACS Synthetic Biology, 2019, 8, 1548-1559.	1.9	27
26	<sc>i</sc>B<sc>io</sc>S<sc>im</sc> 3: A Tool for Model-Based Genetic Circuit Design. ACS Synthetic Biology, 2019, 8, 1560-1563.	1.9	62
27	Specifications of Standards in Systems and Synthetic Biology: Status and Developments in 2019. Journal of Integrative Bioinformatics, 2019, 16, .	1.0	7
28	Specifying Combinatorial Designs with the Synthetic Biology Open Language (SBOL). ACS Synthetic Biology, 2019, 8, 1519-1523.	1.9	2
29	Design of Asynchronous Genetic Circuits. Proceedings of the IEEE, 2019, 107, 1356-1368.	16.4	16
30	Communicating Structure and Function in Synthetic Biology Diagrams. ACS Synthetic Biology, 2019, 8, 1818-1825.	1.9	30
31	Synthetic Biology Open Language Visual (SBOL Visual) Version 2.1. Journal of Integrative Bioinformatics, 2019, 16, .	1.0	8
32	SBOLExplorer: Data Infrastructure and Data Mining for Genetic Design Repositories. ACS Synthetic Biology, 2019, 8, 2287-2294.	1.9	2
33	Synthetic Biology Open Language (SBOL) Version 2.3. Journal of Integrative Bioinformatics, 2019, 16, .	1.0	16
34	The Systems Biology Markup Language (SBML): Language Specification for Level 3 Version 2 Core Release 2. Journal of Integrative Bioinformatics, 2019, 16, .	1.0	78
35	SBOL-OWL: An Ontological Approach for Formal and Semantic Representation of Synthetic Biology Information. ACS Synthetic Biology, 2019, 8, 1498-1514.	1.9	12
36	Harmonizing semantic annotations for computational models in biology. Briefings in Bioinformatics, 2019, 20, 540-550.	3.2	52

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37	sboljs: Bringing the Synthetic Biology Open Language to the Web Browser. ACS Synthetic Biology, 2019, 8, 191-193.	1.9	3
38	pySBOL: A Python Package for Genetic Design Automation and Standardization. ACS Synthetic Biology, 2019, 8, 1515-1518.	1.9	14
39	Toward reproducible disease models using the Systems Biology Markup Language. Simulation, 2019, 95, 895-930.	1.1	3
40	Approximation Techniques for Stochastic Analysis of Biological Systems. Computational Biology, 2019, , 327-348.	0.1	5
41	STAMINA: STochastic Approximate Model-Checker for INfinite-State Analysis. Lecture Notes in Computer Science, 2019, , 540-549.	1.0	9
42	SynBioHub: A Standards-Enabled Design Repository for Synthetic Biology. ACS Synthetic Biology, 2018, 7, 682-688.	1.9	112
43	Specifications of Standards in Systems and Synthetic Biology: Status and Developments in 2017. Journal of Integrative Bioinformatics, 2018, 15, .	1.0	7
44	Synthetic Biology Open Language (SBOL) Version 2.2.0. Journal of Integrative Bioinformatics, 2018, 15, .	1.0	20
45	The Systems Biology Markup Language (SBML): Language Specification for Level 3 Version 1 Core. Journal of Integrative Bioinformatics, 2018, 15, .	1.0	13
46	The Systems Biology Markup Language (SBML): Language Specification for Level 3 Version 2 Core. Journal of Integrative Bioinformatics, 2018, 15, .	1.0	57
47	Synthetic Biology Open Language Visual (SBOL Visual) Version 2.0. Journal of Integrative Bioinformatics, 2018, 15, .	1.0	21
48	SBOLme: a Repository of SBOL Parts for Metabolic Engineering. ACS Synthetic Biology, 2017, 6, 732-736.	1.9	5
49	SBOLDesigner 2: An Intuitive Tool for Structural Genetic Design. ACS Synthetic Biology, 2017, 6, 1150-1160.	1.9	38
50	A standard-enabled workflow for synthetic biology. Biochemical Society Transactions, 2017, 45, 793-803.	1.6	38
51	A Validator and Converter for the Synthetic Biology Open Language. ACS Synthetic Biology, 2017, 6, 1161-1168.	1.9	5
52	Advances in Formal Methods for the Design of Analog/Mixed-Signal Systems. , 2017, , .		1
53	A brief history of COMBINE. , 2017, , .		6
54	Synthetic Biology Open Language (SBOL) Version 2.1.0. Journal of Integrative Bioinformatics, 2016, 13, .	1.0	11

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55	Toward Community Standards and Software for Whole-Cell Modeling. IEEE Transactions on Biomedical Engineering, 2016, 63, 2007-2014.	2.5	51
56	libSBOLj 2.0: A Java Library to Support SBOL 2.0. IEEE Life Sciences Letters, 2016, , 1-1.	1.2	0
57	Sharing Structure and Function in Biological Design with SBOL 2.0. ACS Synthetic Biology, 2016, 5, 498-506.	1.9	88
58	Guest Editors' Introduction Challenges and Opportunities in Analog/Mixed-Signal CAD. IEEE Design and Test, 2016, 33, 5-6.	1.1	2
59	Specifications of Standards in Systems and Synthetic Biology: Status and Developments in 2016. Journal of Integrative Bioinformatics, 2016, 13, 1-7.	1.0	12
60	Design of Mixed-Signal Systems With Asynchronous Control. IEEE Design and Test, 2016, 33, 44-55.	1.1	2
61	An improved fault-tolerant routing algorithm for a Network-on-Chip derived with formal analysis. Science of Computer Programming, 2016, 118, 24-39.	1.5	9
62	Efficient Analysis of Systems Biology Markup Language Models of Cellular Populations Using Arrays. ACS Synthetic Biology, 2016, 5, 835-841.	1.9	5
63	A Converter from the Systems Biology Markup Language to the Synthetic Biology Open Language. ACS Synthetic Biology, 2016, 5, 479-486.	1.9	20
64	Synthetic Biology Open Language (SBOL) Version 2.1.0. Journal of Integrative Bioinformatics, 2016, 13, 291.	1.0	6
65	Specifications of Standards in Systems and Synthetic Biology: Status and Developments in 2016. Journal of Integrative Bioinformatics, 2016, 13, 289.	1.0	7
66	libSBOLj 2.0: A Java Library to Support SBOL 2.0. IEEE Life Sciences Letters, 2015, 1, 34-37.	1.2	24
67	Specifications of Standards in Systems and Synthetic Biology. Journal of Integrative Bioinformatics, 2015, 12, 1-3.	1.0	11
68	Synthetic Biology Open Language (SBOL) Version 2.0.0. Journal of Integrative Bioinformatics, 2015, 12, 902-991.	1.0	22
69	Promoting Coordinated Development of Community-Based Information Standards for Modeling in Biology: The COMBINE Initiative. Frontiers in Bioengineering and Biotechnology, 2015, 3, 19.	2.0	72
70	SBML Level 3 package: Hierarchical Model Composition, Version 1 Release 3. Journal of Integrative Bioinformatics, 2015, 12, 603-659.	1.0	39
71	Generating Systems Biology Markup Language Models from the Synthetic Biology Open Language. ACS Synthetic Biology, 2015, 4, 873-879.	1.9	81
72	Computational Synthetic Biology: Progress and the Road Ahead. IEEE Transactions on Multi-Scale Computing Systems, 2015, 1, 19-32.	2.5	7

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73	JSBML 1.0: providing a smorgasbord of options to encode systems biology models. <i>Bioinformatics</i> , 2015, 31, 3383-3386.	1.8	37
74	Proposed Data Model for the Next Version of the Synthetic Biology Open Language. <i>ACS Synthetic Biology</i> , 2015, 4, 57-71.	1.9	19
75	The Synthetic Biology Open Language. <i>Methods in Molecular Biology</i> , 2015, 1244, 323-336.	0.4	1
76	SBOL Visual: A Graphical Language for Genetic Designs. <i>PLoS Biology</i> , 2015, 13, e1002310.	2.6	73
77	Specifications of Standards in Systems and Synthetic Biology. <i>Journal of Integrative Bioinformatics</i> , 2015, 12, 258.	1.0	14
78	SBML Level 3 package: Hierarchical Model Composition, Version 1 Release 3. <i>Journal of Integrative Bioinformatics</i> , 2015, 12, 268.	1.0	31
79	Systems Biology Markup Language (SBML) Level 2 Version 5: Structures and Facilities for Model Definitions. <i>Journal of Integrative Bioinformatics</i> , 2015, 12, 271.	1.0	42
80	Synthetic Biology Open Language (SBOL) Version 2.0.0. <i>Journal of Integrative Bioinformatics</i> , 2015, 12, 272.	1.0	21
81	Efficient Analysis Methods in Synthetic Biology. <i>Methods in Molecular Biology</i> , 2015, 1244, 217-257.	0.4	0
82	Hierarchical Stochastic Simulation Algorithm for SBML Models of Genetic Circuits. <i>Frontiers in Bioengineering and Biotechnology</i> , 2014, 2, 55.	2.0	5
83	Stochastic Model Checking of Genetic Circuits. <i>ACM Journal on Emerging Technologies in Computing Systems</i> , 2014, 11, 1-21.	1.8	10
84	Introduction to the Special Issue on Computational Synthetic Biology. <i>ACM Journal on Emerging Technologies in Computing Systems</i> , 2014, 11, 1-5.	1.8	0
85	LEMA: A tool for the formal verification of digitally-intensive analog/mixed-signal circuits. , 2014, , .		2
86	A Methodology to Annotate Systems Biology Markup Language Models with the Synthetic Biology Open Language. <i>ACS Synthetic Biology</i> , 2014, 3, 57-66.	1.9	30
87	Directed Acyclic Graph-Based Technology Mapping of Genetic Circuit Models. <i>ACS Synthetic Biology</i> , 2014, 3, 543-555.	1.9	30
88	Compositional Model Checking of Concurrent Systems. <i>IEEE Transactions on Computers</i> , 2014, , 1-1.	2.4	3
89	The Synthetic Biology Open Language (SBOL) provides a community standard for communicating designs in synthetic biology. <i>Nature Biotechnology</i> , 2014, 32, 545-550.	9.4	247
90	Meeting report from the fourth meeting of the Computational Modeling in Biology Network (COMBINE). <i>Standards in Genomic Sciences</i> , 2014, 9, 1285-1301.	1.5	21

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91	Formal Analysis of a Fault-Tolerant Routing Algorithm for a Network-on-Chip. Lecture Notes in Computer Science, 2014, , 48-62.	1.0	3
92	Verification of digitally-intensive analog circuits via kernel ridge regression and hybrid reachability analysis. , 2013, , .		9
93	Dynamic Modeling of Cellular Populations within iBioSim. ACS Synthetic Biology, 2013, 2, 223-229.	1.9	15
94	IWBDA 2012 Special Issue. ACS Synthetic Biology, 2013, 2, 203-203.	1.9	0
95	Platforms for Genetic Design Automation. Methods in Microbiology, 2013, , 177-202.	0.4	6
96	An Improvement in Partial Order Reduction Using Behavioral Analysis. , 2012, , .		2
97	Using decision diagrams to compactly represent the state space for explicit model checking. , 2012, , .		2
98	Utilizing stochastic model checking to analyze genetic circuits. , 2012, , .		11
99	Poster Abstract: Methods and Tools for Verification of Cyber-Physical Systems. , 2012, , .		0
100	<title></title> </titles> </publication_date> </month>06</month> </year>2012</year> </publication_date> </pages> </first_page>C1</first_page> </last_page>C1</last_page> </pages> </publisher_item> </item_number item_number_type='arNumber'>6327710</item_number> </publisher_item> </doi_data> </doi>10.1109/MDT.2012.2211731</doi> </resource>http://ieeexplore.ieee.org/lpdocs/epic03/wrapper.htm?arnumbe.	1.4	31
101	A Compositional Minimization Approach for Large Asynchronous Design Verification. Lecture Notes in Computer Science, 2012, , 62-79.	1.0	2
102	Formal Verification of Genetic Circuits. Lecture Notes in Computer Science, 2012, , 5-5.	1.0	1
103	Learning Genetic Regulatory Network Connectivity from Time Series Data. IEEE/ACM Transactions on Computational Biology and Bioinformatics, 2011, 8, 152-165.	1.9	19
104	A Behavioral Analysis Approach for Efficient Partial Order Reduction. , 2011, , .		1
105	Verification of Analog/Mixed-Signal Circuits Using Labeled Hybrid Petri Nets. IEEE Transactions on Computer-Aided Design of Integrated Circuits and Systems, 2011, 30, 617-630.	1.9	17
106	Erlang-delayed stochastic chemical kinetic formalism for efficient analysis of biological systems with non-elementary reaction effects. , 2011, , .		0
107	Design and analysis of a robust genetic Muller C-element. Journal of Theoretical Biology, 2010, 264, 174-187.	0.8	12
108	iSSA: An incremental stochastic simulation algorithm for genetic circuits. , 2010, , .		5

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109	ANALOG/MIXED-SIGNAL CIRCUIT VERIFICATION USING MODELS GENERATED FROM SIMULATION TRACES. International Journal of Foundations of Computer Science, 2010, 21, 191-210.	0.8	22
110	Temperature Control of Fimbriation Circuit Switch in Uropathogenic Escherichia coli: Quantitative Analysis via Automated Model Abstraction. PLoS Computational Biology, 2010, 6, e1000723.	1.5	21
111	State space reductions for scalable verification of asynchronous designs. , 2010, , .		0
112	Automatic abstraction for verification of cyber-physical systems. , 2010, , .		45
113	Abstraction Methods for Analysis of Gene Regulatory Networks. , 2010, , 352-385.		0
114	iBioSim: a tool for the analysis and design of genetic circuits. Bioinformatics, 2009, 25, 2848-2849.	1.8	100
115	A new verification method for embedded systems. , 2009, , .		5
116	A Behavioral Synthesis System for Asynchronous Circuits with Bundled-data Implementation. IPSJ Transactions on System LSI Design Methodology, 2009, 2, 64-79.	0.5	3
117	Verification of Analog/Mixed-Signal Circuits Using Symbolic Methods. IEEE Transactions on Computer-Aided Design of Integrated Circuits and Systems, 2008, 27, 2223-2235.	1.9	23
118	A behavioral synthesis method for asynchronous circuits with bundled-data implementation (Tool) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50		
119	Production-Passage-Time Approximation: A New Approximation Method to Accelerate the Simulation Process of Enzymatic Reactions. Journal of Computational Biology, 2008, 15, 779-792.	0.8	5
120	A Conservative Framework for Safety-Failure Checking. IEICE Transactions on Information and Systems, 2008, E91-D, 642-654.	0.4	0
121	The Design of a Genetic Muller C-Element. Proceedings of the International Symposium on Advanced Research in Asynchronous Circuits and Systems, 2007, , .	0.0	14
122	Efficient Verification of Hazard-Freedom in Gate-Level Timed Asynchronous Circuits. IEEE Transactions on Computer-Aided Design of Integrated Circuits and Systems, 2007, 26, 592-605.	1.9	3
123	Synthesis of Timed Circuits Based on Decomposition. IEEE Transactions on Computer-Aided Design of Integrated Circuits and Systems, 2007, 26, 1177-1195.	1.9	5
124	Application of Automated Model Generation Techniques to Analog/Mixed-Signal Circuits. , 2007, , .		2
125	Hazard Checking of Timed Asynchronous Circuits Revisited. International Conference on Application of Concurrency To System Design, 2007, , .	0.0	0
126	Symbolic Model Checking of Analog/Mixed-Signal Circuits. , 2007, , .		6

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127	Analog/Mixed-Signal Circuit Verification Using Models Generated from Simulation Traces. , 2007, , 114-128.		18
128	Bounded Model Checking of Analog and Mixed-Signal Circuits Using an SMT Solver. , 2007, , 66-81.		5
129	Scheduling Methods for Asynchronous Circuits with Bundled-Data Implementations Based on the Approximation of Start Times. IEICE Transactions on Fundamentals of Electronics, Communications and Computer Sciences, 2007, E90-A, 2790-2799.	0.2	9
130	Verification of Analog/Mixed-Signal Circuits Using Labeled Hybrid Petri Nets. IEEE/ACM International Conference on Computer-Aided Design, Digest of Technical Papers, 2006, , .	0.0	14
131	Verification of timed circuits with failure-directed abstractions. IEEE Transactions on Computer-Aided Design of Integrated Circuits and Systems, 2006, 25, 403-412.	1.9	10
132	The Case for Analog Circuit Verification. Electronic Notes in Theoretical Computer Science, 2006, 153, 53-63.	0.9	15
133	Verification of analog/mixed-signal circuits using labeled hybrid petri nets. IEEE/ACM International Conference on Computer-Aided Design, Digest of Technical Papers, 2006, , .	0.0	17
134	ILP-based Scheduling for Asynchronous Circuits in Bundled-Data Implementation. , 2006, , .		4
135	Effective Contraction of Timed STGs for Decomposition Based Timed Circuit Synthesis. Lecture Notes in Computer Science, 2006, , 229-244.	1.0	0
136	Partial Order Reduction for Detecting Safety and Timing Failures of Timed Circuits. IEICE Transactions on Information and Systems, 2005, E88-D, 1646-1661.	0.4	7
137	Failure Trace Analysis of Timed Circuits for Automatic Timing Constraints Derivation. IEICE Transactions on Information and Systems, 2005, E88-D, 2555-2564.	0.4	0
138	CMOS Analog MAP Decoder for (8,4) Hamming Code. IEEE Journal of Solid-State Circuits, 2004, 39, 122-131.	3.5	64
139	Verification of Analog and Mixed-Signal Circuits Using Timed Hybrid Petri Nets. Lecture Notes in Computer Science, 2004, , 426-440.	1.0	12
140	Modular verification of timed circuits using automatic abstraction. IEEE Transactions on Computer-Aided Design of Integrated Circuits and Systems, 2003, 22, 1138-1153.	1.9	19
141	Direct synthesis of timed circuits from free-choice STGs. IEEE Transactions on Computer-Aided Design of Integrated Circuits and Systems, 2002, 21, 275-290.	1.9	1
142	Efficient algorithms for exact two-level hazard-free logic minimization. IEEE Transactions on Computer-Aided Design of Integrated Circuits and Systems, 2002, 21, 1269-1283.	1.9	12
143	Modular Synthesis of Timed Circuits using Partial Orders on LPNs. Electronic Notes in Theoretical Computer Science, 2002, 65, 180-201.	0.9	2
144	An asynchronous instruction length decoder. IEEE Journal of Solid-State Circuits, 2001, 36, 217-228.	3.5	55

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145	Timed circuit verification using TEL structures. IEEE Transactions on Computer-Aided Design of Integrated Circuits and Systems, 2001, 20, 129-146.	1.9	12
146	Timed circuits. , 2001, , .		20
147	Automatic Abstraction for Verification of Timed Circuits and Systems?. Lecture Notes in Computer Science, 2001, , 182-193.	1.0	5
148	Interfacing synchronous and asynchronous modules within a high-speed pipeline. IEEE Transactions on Very Large Scale Integration (VLSI) Systems, 2000, 8, 573-583.	2.1	33
149	Timed state space exploration using POSETs. IEEE Transactions on Computer-Aided Design of Integrated Circuits and Systems, 2000, 19, 501-520.	1.9	21
150	POSET timing and its application to the synthesis and verification of gate-level timed circuits. IEEE Transactions on Computer-Aided Design of Integrated Circuits and Systems, 1999, 18, 769-786.	1.9	13
151	Covering conditions and algorithms for the synthesis of speed-independent circuits. IEEE Transactions on Computer-Aided Design of Integrated Circuits and Systems, 1998, 17, 205-219.	1.9	21
152	Verification of timed systems using POSETs. Lecture Notes in Computer Science, 1998, , 403-415.	1.0	29
153	Synthesis of timed asynchronous circuits. IEEE Transactions on Very Large Scale Integration (VLSI) Systems, 1993, 1, 106-119.	2.1	92
154	Engineering Genetic Circuits. , 0, , .		16
155	Efficient Stochastic Simulation to Analyze Targeted Properties of Biological Systems. , 0, , .		3