## Sarah M Keating

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4002834/publications.pdf

Version: 2024-02-01

30 2,530 14 28 papers citations h-index g-index

32 32 32 32 3804

times ranked

citing authors

docs citations

all docs

#	Article	IF	Citations
1	Creation and analysis of biochemical constraint-based models using the COBRA Toolbox v.3.0. Nature Protocols, 2019, 14, 639-702.	12.0	833
2	LibSBML: an API Library for SBML. Bioinformatics, 2008, 24, 880-881.	4.1	342
3	Evolving a lingua franca and associated software infrastructure for computational systems biology: the Systems Biology Markup Language (SBML) project. IET Systems Biology, 2004, 1, 41-53.	2.0	187
4	<scp>SBML</scp> Level 3: an extensible format for the exchange and reuse of biological models. Molecular Systems Biology, 2020, 16, e9110.	7.2	178
5	BioModels—15 years of sharing computational models in life science. Nucleic Acids Research, 2020, 48, D407-D415.	14.5	175
6	SBML qualitative models: a model representation format and infrastructure to foster interactions between qualitative modelling formalisms and tools. BMC Systems Biology, 2013, 7, 135.	3.0	145
7	SBMLToolbox: an SBML toolbox for MATLAB users. Bioinformatics, 2006, 22, 1275-1277.	4.1	129
8	The Systems Biology Markup Language (SBML): Language Specification for Level 3 Version 1 Core. Journal of Integrative Bioinformatics, 2015, 12, 266.	1.5	102
9	The Systems Biology Markup Language (SBML): Language Specification for Level 3 Version 2 Core Release 2. Journal of Integrative Bioinformatics, 2019, 16, .	1.5	78
10	Wikidata as a knowledge graph for the life sciences. ELife, 2020, 9, .	6.0	76
11	The Systems Biology Markup Language (SBML): Language Specification for Level 3 Version 2 Core. Journal of Integrative Bioinformatics, 2018, 15, .	1.5	57
12	Systems Biology Markup Language (SBML) Level 2 Version 5: Structures and Facilities for Model Definitions. Journal of Integrative Bioinformatics, 2015, 12, 271.	1.5	42
13	The Systems Biology Markup Language (SBML) Level 3 Package: Qualitative Models, Version 1, Release 1. Journal of Integrative Bioinformatics, 2015, 12, 270.	1.5	21
14	SBML Level 3 package: Render, Version 1, Release 1. Journal of Integrative Bioinformatics, 2018, 15, .	1.5	18
15	The first 10 years of the international coordination network for standards in systems and synthetic biology (COMBINE). Journal of Integrative Bioinformatics, 2020, 17, .	1.5	18
16	SBML Level 3 package: Qualitative Models, Version 1, Release 1. Journal of Integrative Bioinformatics, 2015, 12, 691-730.	1.5	15
17	MOCCASIN: converting MATLAB ODE models to SBML. Bioinformatics, 2016, 32, 1905-1906.	4.1	14
18	The Systems Biology Markup Language (SBML): Language Specification for Level 3 Version 1 Core. Journal of Integrative Bioinformatics, 2018, 15, .	1.5	13

#	Article	IF	CITATIONS
19	Supporting SBML as a Model Exchange Format in Software Applications. Methods in Molecular Biology, 2013, 1021, 201-225.	0.9	12
20	Methods for Simulating the Dynamics of Complex Biological Processes. Methods in Cell Biology, 2008, 84, 807-842.	1.1	10
21	Specifications of standards in systems and synthetic biology: status and developments in 2020. Journal of Integrative Bioinformatics, 2020, 17, .	1.5	10
22	Addressing <i>barriers in comprehensiveness, accessibility, reusability, interoperability and reproducibility of computational models in systems biology</i>  i>. Briefings in Bioinformatics, 2022, 23, .	6.5	10
23	A Profile of Today's SBML-Compatible Software. , 2011, , .		9
24	Systems biology markup language (SBML) level 3 package: multistate, multicomponent and multicompartment species, version 1, release 2. Journal of Integrative Bioinformatics, 2020, 17, .	1.5	8
25	Specifications of Standards in Systems and Synthetic Biology: Status and Developments in 2017. Journal of Integrative Bioinformatics, 2018, 15, .	1.5	7
26	Specifications of Standards in Systems and Synthetic Biology: Status and Developments in 2019. Journal of Integrative Bioinformatics, 2019, 16, .	1.5	7
27	Systems Biology Markup Language (SBML) Level 3 Package: Distributions, Version 1, Release 1. Journal of Integrative Bioinformatics, 2020, 17, .	1.5	7
28	SBML Models and MathSBML., 2007,, 395-421.		3
29	Specifications of standards in systems and synthetic biology: status and developments in 2021. Journal of Integrative Bioinformatics, 2021, 18, .	1.5	2
30	chaste codegen: automatic CellML to C++ code generation with fixes for singularities and automatically generated Jacobians. Wellcome Open Research, 2021, 6, 261.	1.8	1