

# Allyson L Lister

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6081571/publications.pdf>

Version: 2024-02-01

17  
papers

3,016  
citations

759055

12  
h-index

794469

19  
g-index

21  
all docs

21  
docs citations

21  
times ranked

5744  
citing authors

#	ARTICLE	IF	CITATIONS
1	Helping the Consumers and Producers of Standards, Repositories and Policies to Enable FAIR Data. Data Intelligence, 2020, 2, 151-157.	0.8	10
2	<scp>SBML</scp> Level 3: an extensible format for the exchange and reuse of biological models. Molecular Systems Biology, 2020, 16, e9110.	3.2	178
3	FAIRsharing as a community approach to standards, repositories and policies. Nature Biotechnology, 2019, 37, 358-367.	9.4	228
4	The Ontology for Biomedical Investigations. PLoS ONE, 2016, 11, e0154556.	1.1	217
5	The Software Ontology (SWO): a resource for reproducibility in biomedical data analysis, curation and digital preservation. Journal of Biomedical Semantics, 2014, 5, 25.	0.9	56
6	MIREOT: The minimum information to reference an external ontology term. Applied Ontology, 2011, 6, 23-33.	1.0	78
7	Controlled vocabularies and semantics in systems biology. Molecular Systems Biology, 2011, 7, 543.	3.2	246
8	Quantitative Fitness Analysis Shows That NMD Proteins and Many Other Protein Complexes Suppress or Enhance Distinct Telomere Cap Defects. PLoS Genetics, 2011, 7, e1001362.	1.5	65
9	Annotation of SBML models through rule-based semantic integration. Journal of Biomedical Semantics, 2010, 1, S3.	0.9	16
10	Live Coverage of Scientific Conferences Using Web Technologies. PLoS Computational Biology, 2010, 6, e1000563.	1.5	8
11	Live Coverage of Intelligent Systems for Molecular Biology/European Conference on Computational Biology (ISMB/ECCB) 2009. PLoS Computational Biology, 2010, 6, e1000640.	1.5	4
12	Managing Experimental Data Using FuGE. Methods in Molecular Biology, 2010, 604, 333-343.	0.4	4
13	Modeling and Managing Experimental Data Using FuGE. OMICS A Journal of Integrative Biology, 2009, 13, 239-251.	1.0	8
14	Saint: a lightweight integration environment for model annotation. Bioinformatics, 2009, 25, 3026-3027.	1.8	25
15	The minimum information about a genome sequence (MIGS) specification. Nature Biotechnology, 2008, 26, 541-547.	9.4	1,069
16	eGenomics: Cataloguing Our Complete Genome Collection III. Comparative and Functional Genomics, 2007, 2007, 1-7.	2.0	4
17	The International Protein Index: An integrated database for proteomics experiments. Proteomics, 2004, 4, 1985-1988.	1.3	685