Allyson L Lister

List of Publications by Year
in descending order

17 papers	3,016 citations	759233 12 h-index	794594 19 g-index
21 all docs	21 docs citations	21 times ranked	5744 citing authors

#	Article	IF	Citations
1	The minimum information about a genome sequence (MIGS) specification. Nature Biotechnology, 2008, 26, 541-547.	17.5	1,069
2	The International Protein Index: An integrated database for proteomics experiments. Proteomics, 2004, 4, 1985-1988.	2.2	685
3	Controlled vocabularies and semantics in systems biology. Molecular Systems Biology, 2011, 7, 543.	7.2	246
4	FAIRsharing as a community approach to standards, repositories and policies. Nature Biotechnology, 2019, 37, 358-367.	17.5	228
5	The Ontology for Biomedical Investigations. PLoS ONE, 2016, 11, e0154556.	2.5	217
6	<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
7	MIREOT: The minimum information to reference an external ontology term. Applied Ontology, 2011, 6, 23-33.	2.0	78
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.	3.5	65
9	The Software Ontology (SWO): a resource for reproducibility in biomedical data analysis, curation and digital preservation. Journal of Biomedical Semantics, 2014, 5, 25.	1.6	56
10	Saint: a lightweight integration environment for model annotation. Bioinformatics, 2009, 25, 3026-3027.	4.1	25
11	Annotation of SBML models through rule-based semantic integration. Journal of Biomedical Semantics, 2010, 1, S3.	1.6	16
12	Helping the Consumers and Producers of Standards, Repositories and Policies to Enable FAIR Data. Data Intelligence, 2020, 2, 151-157.	1.5	10
13	Modeling and Managing Experimental Data Using FuGE. OMICS A Journal of Integrative Biology, 2009, 13, 239-251.	2.0	8
14	Live Coverage of Scientific Conferences Using Web Technologies. PLoS Computational Biology, 2010, 6, e1000563.	3.2	8
15	eGenomics: Cataloguing Our Complete Genome Collection III. Comparative and Functional Genomics, 2007, 2007, 1-7.	2.0	4
16	Live Coverage of Intelligent Systems for Molecular Biology/European Conference on Computational Biology (ISMB/ECCB) 2009. PLoS Computational Biology, 2010, 6, e1000640.	3.2	4
17	Managing Experimental Data Using FuGE. Methods in Molecular Biology, 2010, 604, 333-343.	0.9	4