

# CITATION REPORT

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## Curation accuracy of model organism databases

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Database: the Journal of Biological Databases and Curation, 2014, 2014, .

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#	Paper	IF	Citations
25	The Comparative Toxicogenomics Database's 10th year anniversary: update 2015. <i>Nucleic Acids Research</i> , <b>2015</b> , 43, D914-20	20.1	288
24	How much does curation cost?. <i>Database: the Journal of Biological Databases and Curation</i> , <b>2016</b> , 2016,	5	35
23	Identification of Unannotated Small Genes in. <i>G3: Genes, Genomes, Genetics</i> , <b>2017</b> , 7, 983-989	3.2	39
22	On expert curation and scalability: UniProtKB/Swiss-Prot as a case study. <i>Bioinformatics</i> , <b>2017</b> , 33, 3454-3460	3.6	60
21	A statistical approach to identify, monitor, and manage incomplete curated data sets. <i>BMC Bioinformatics</i> , <b>2018</b> , 19, 110	3.6	1
20	Crowd enabled curation and querying of large and noisy text mined protein interaction data. <i>Distributed and Parallel Databases</i> , <b>2018</b> , 36, 9-45	0.9	2
19	The EcoCyc Database. <i>EcoSal Plus</i> , <b>2018</b> , 8,	7.7	29
18	Scaling up data curation using deep learning: An application to literature triage in genomic variation resources. <i>PLoS Computational Biology</i> , <b>2018</b> , 14, e1006390	5	21
17	SamPler - a novel method for selecting parameters for gene functional annotation routines. <i>BMC Bioinformatics</i> , <b>2019</b> , 20, 454	3.6	3
16	The BioCyc collection of microbial genomes and metabolic pathways. <i>Briefings in Bioinformatics</i> , <b>2019</b> , 20, 1085-1093	13.4	232
15	Detecting Gene Ontology misannotations using taxon-specific rate ratio comparisons. <i>Bioinformatics</i> , <b>2020</b> , 36, 4383-4388	7.2	6
14	Constructing knowledge graphs and their biomedical applications. <i>Computational and Structural Biotechnology Journal</i> , <b>2020</b> , 18, 1414-1428	6.8	36
13	Metabolic networks of the Nicotiana genus in the spotlight: content, progress and outlook. <i>Briefings in Bioinformatics</i> , <b>2021</b> , 22,	13.4	0
12	Leveraging Curation Among Pathway/Genome Databases Using Ortholog-Based Annotation Propagation. <i>Frontiers in Microbiology</i> , <b>2021</b> , 12, 614355	5.7	1
11	Can we replace curation with information extraction software?. <i>Database: the Journal of Biological Databases and Curation</i> , <b>2016</b> , 2016,	5	13
10	Towards Knowledge Maintenance in Scientific Digital Libraries with the Keystone Framework. <b>2020</b> , 2020, 217-226		3
9	Funding knowledgebases: Towards a sustainable funding model for the UniProt use case. <i>F1000Research</i> , <b>2017</b> , 6,	3.6	12

8	Funding knowledgebases: Towards a sustainable funding model for the UniProt use case. <i>F1000Research</i> , <b>2017</b> , 6, 2051	3.6	13
7	Insights from the reanalysis of high-throughput chemical genomics data for Escherichia coli K-12. <i>G3: Genes, Genomes, Genetics</i> , <b>2021</b> , 11,	3.2	0
6	Modelling Oxidative Stress Pathways. <i>Computational Biology</i> , <b>2020</b> , 277-300	0.7	
5	Insights from the reanalysis of high-throughput chemical genomics data for Escherichia coli K-12.		
4	Undergraduate Biocuration: Developing Tomorrow's Researchers While Mining Today's Data. <i>Journal of Undergraduate Neuroscience Education: JUNE: A Publication of FUN, Faculty for Undergraduate Neuroscience</i> , <b>2015</b> , 14, A56-65	0.6	19
3	Evaluating the predictive accuracy of curated biological pathways in a public knowledgebase.. <i>Database: the Journal of Biological Databases and Curation</i> , <b>2022</b> , 2022,	5	0
2	FoodChem: A food-chemical relation extraction model. <b>2021</b> ,		
1	Ecosystem-specific microbiota and microbiome databases in the era of big data. <i>Environmental Microbiomes</i> , <b>2022</b> , 17,	5.6	1