

Rebecca E Foulger

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1764734/publications.pdf>

Version: 2024-02-01

10
papers

1,227
citations

933447

10
h-index

1372567

10
g-index

11
all docs

11
docs citations

11
times ranked

2590
citing authors

#	ARTICLE	IF	CITATIONS
1	Scaling national and international improvement in virtual gene panel curation via a collaborative approach to discordance resolution. <i>American Journal of Human Genetics</i> , 2021, 108, 1551-1557.	6.2	36
2	SynGO: An Evidence-Based, Expert-Curated Knowledge Base for the Synapse. <i>Neuron</i> , 2019, 103, 217-234.e4.	8.1	518
3	Improving Interpretation of Cardiac Phenotypes and Enhancing Discovery With Expanded Knowledge in the Gene Ontology. <i>Circulation Genomic and Precision Medicine</i> , 2018, 11, e001813.	3.6	24
4	An expanded evaluation of protein function prediction methods shows an improvement in accuracy. <i>Genome Biology</i> , 2016, 17, 184.	8.8	308
5	Representing Kidney Development Using the Gene Ontology. <i>PLoS ONE</i> , 2014, 9, e99864.	2.5	17
6	TermGenie – a web-application for pattern-based ontology class generation. <i>Journal of Biomedical Semantics</i> , 2014, 5, 48.	1.6	30
7	A method for increasing expressivity of Gene Ontology annotations using a compositional approach. <i>BMC Bioinformatics</i> , 2014, 15, 155.	2.6	78
8	The Gene Ontology (GO) Cellular Component Ontology: integration with SAO (Subcellular Anatomy) Tj ETQq0 0 0 rgBT /Overlock 10 Tf	1.6	44
9	Dovetailing biology and chemistry: integrating the Gene Ontology with the ChEBI chemical ontology. <i>BMC Genomics</i> , 2013, 14, 513.	2.8	45
10	Predicting Gene Function From Patterns of Annotation. <i>Genome Research</i> , 2003, 13, 896-904.	5.5	121