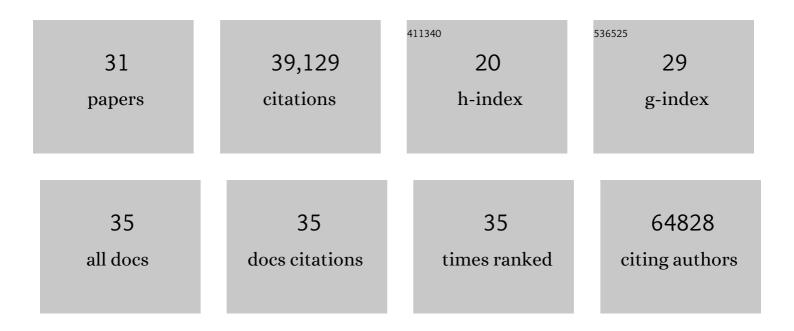
Midori A Harris

List of Publications by Year in descending order

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ΜΙΠΟΡΙ Δ ΗΛΟΡΙς

#	Article	IF	CITATIONS
1	JaponicusDB: rapid deployment of a model organism database for an emerging model species. Genetics, 2022, 220, .	1.2	21
2	Fission stories: using PomBase to understand <i>Schizosaccharomyces pombe</i> biology. Genetics, 2022, 220, .	1.2	60
3	The Gene Ontology resource: enriching a GOld mine. Nucleic Acids Research, 2021, 49, D325-D334.	6.5	2,416
4	Community curation in PomBase: enabling fission yeast experts to provide detailed, standardized, sharable annotation from research publications. Database: the Journal of Biological Databases and Curation, 2020, 2020, .	1.4	19
5	Term Matrix: a novel Gene Ontology annotation quality control system based on ontology term co-annotation patterns. Open Biology, 2020, 10, 200149.	1.5	7
6	PomBase 2018: user-driven reimplementation of the fission yeast database provides rapid and intuitive access to diverse, interconnected information. Nucleic Acids Research, 2019, 47, D821-D827.	6.5	157
7	Hidden in plain sight: what remains to be discovered in the eukaryotic proteome?. Open Biology, 2019, 9, 180241.	1.5	80
8	Annotation of gene product function from high-throughput studies using the Gene Ontology. Database: the Journal of Biological Databases and Curation, 2019, 2019, .	1.4	21
9	PomBase: The Scientific Resource for Fission Yeast. Methods in Molecular Biology, 2018, 1757, 49-68.	0.4	32
10	Model organism databases: essential resources that need the support of both funders and users. BMC Biology, 2016, 14, 49.	1.7	46
11	PomBase 2015: updates to the fission yeast database. Nucleic Acids Research, 2015, 43, D656-D661.	6.5	95
12	Representing Kidney Development Using the Gene Ontology. PLoS ONE, 2014, 9, e99864.	1.1	17
13	Canto: an online tool for community literature curation. Bioinformatics, 2014, 30, 1791-1792.	1.8	41
14	A method for increasing expressivity of Gene Ontology annotations using a compositional approach. BMC Bioinformatics, 2014, 15, 155.	1.2	78
15	Dovetailing biology and chemistry: integrating the Gene Ontology with the ChEBI chemical ontology. BMC Genomics, 2013, 14, 513.	1.2	45
16	A guide to best practices for Gene Ontology (GO) manual annotation. Database: the Journal of Biological Databases and Curation, 2013, 2013, bat054-bat054.	1.4	135
17	FYPO: the fission yeast phenotype ontology. Bioinformatics, 2013, 29, 1671-1678.	1.8	53
18	Semantic integration of physiology phenotypes with an application to the Cellular Phenotype Ontology. Bioinformatics, 2012, 28, 1783-1789.	1.8	22

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#	Article	IF	CITATIONS
19	PomBase: a comprehensive online resource for fission yeast. Nucleic Acids Research, 2012, 40, D695-D699.	6.5	288
20	Cross-product extensions of the Gene Ontology. Journal of Biomedical Informatics, 2011, 44, 80-86.	2.5	96
21	How the gene ontology evolves. BMC Bioinformatics, 2011, 12, 325.	1.2	32
22	Ontology engineering. Nature Biotechnology, 2010, 28, 128-130.	9.4	113
23	The Protein Feature Ontology: a tool for the unification of protein feature annotations. Bioinformatics, 2008, 24, 2767-2772.	1.8	19
24	Developing an Ontology. Methods in Molecular Biology, 2008, 452, 111-124.	0.4	9
25	OBO-Edit an ontology editor for biologists. Bioinformatics, 2007, 23, 2198-2200.	1.8	250
26	Standards and Ontologies for Functional Genomics 2. Comparative and Functional Genomics, 2004, 5, 618-622.	2.0	6
27	Standards and Ontologies for Functional Genomics: Towards Unified Ontologies for Biology and Biomedicine. Comparative and Functional Genomics, 2003, 4, 116-120.	2.0	4
28	The Gene Ontology (GO) Project: Structured Vocabularies for Molecular Biology and Their Application to Genome and Expression Analysis. Current Protocols in Bioinformatics, 2003, 00, Unit 7.2.	25.8	23
29	Gene Ontology: tool for the unification of biology. Nature Genetics, 2000, 25, 25-29.	9.4	34,499
30	Expanding yeast knowledge online. , 1998, 14, 1453-1469.		11
31	Comparison of the Complete Protein Sets of Worm and Yeast: Orthology and Divergence. , 1998, 282, 2022-2028.		404