

# Midori A Harris

## List of Publications by Year in descending order

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Version: 2024-02-01

31  
papers

39,129  
citations

361045

20  
h-index

476904

29  
g-index

35  
all docs

35  
docs citations

35  
times ranked

58454  
citing authors

#	ARTICLE	IF	CITATIONS
1	Gene Ontology: tool for the unification of biology. <i>Nature Genetics</i> , 2000, 25, 25-29.	9.4	34,499
2	The Gene Ontology resource: enriching a GOld mine. <i>Nucleic Acids Research</i> , 2021, 49, D325-D334.	6.5	2,416
3	Comparison of the Complete Protein Sets of Worm and Yeast: Orthology and Divergence. , 1998, 282, 2022-2028.		404
4	PomBase: a comprehensive online resource for fission yeast. <i>Nucleic Acids Research</i> , 2012, 40, D695-D699.	6.5	288
5	OBO-Edit an ontology editor for biologists. <i>Bioinformatics</i> , 2007, 23, 2198-2200.	1.8	250
6	PomBase 2018: user-driven reimplementaion of the fission yeast database provides rapid and intuitive access to diverse, interconnected information. <i>Nucleic Acids Research</i> , 2019, 47, D821-D827.	6.5	157
7	A guide to best practices for Gene Ontology (GO) manual annotation. <i>Database: the Journal of Biological Databases and Curation</i> , 2013, 2013, bat054-bat054.	1.4	135
8	Ontology engineering. <i>Nature Biotechnology</i> , 2010, 28, 128-130.	9.4	113
9	Cross-product extensions of the Gene Ontology. <i>Journal of Biomedical Informatics</i> , 2011, 44, 80-86.	2.5	96
10	PomBase 2015: updates to the fission yeast database. <i>Nucleic Acids Research</i> , 2015, 43, D656-D661.	6.5	95
11	Hidden in plain sight: what remains to be discovered in the eukaryotic proteome?. <i>Open Biology</i> , 2019, 9, 180241.	1.5	80
12	A method for increasing expressivity of Gene Ontology annotations using a compositional approach. <i>BMC Bioinformatics</i> , 2014, 15, 155.	1.2	78
13	Fission stories: using PomBase to understand <i>Schizosaccharomyces pombe</i> biology. <i>Genetics</i> , 2022, 220, .	1.2	60
14	FYPO: the fission yeast phenotype ontology. <i>Bioinformatics</i> , 2013, 29, 1671-1678.	1.8	53
15	Model organism databases: essential resources that need the support of both funders and users. <i>BMC Biology</i> , 2016, 14, 49.	1.7	46
16	Dovetailing biology and chemistry: integrating the Gene Ontology with the ChEBI chemical ontology. <i>BMC Genomics</i> , 2013, 14, 513.	1.2	45
17	Canto: an online tool for community literature curation. <i>Bioinformatics</i> , 2014, 30, 1791-1792.	1.8	41
18	How the gene ontology evolves. <i>BMC Bioinformatics</i> , 2011, 12, 325.	1.2	32

#	ARTICLE	IF	CITATIONS
19	PomBase: The Scientific Resource for Fission Yeast. <i>Methods in Molecular Biology</i> , 2018, 1757, 49-68.	0.4	32
20	The Gene Ontology ( GO ) Project: Structured Vocabularies for Molecular Biology and Their Application to Genome and Expression Analysis. <i>Current Protocols in Bioinformatics</i> , 2003, 00, Unit 7.2.	25.8	23
21	Semantic integration of physiology phenotypes with an application to the Cellular Phenotype Ontology. <i>Bioinformatics</i> , 2012, 28, 1783-1789.	1.8	22
22	Annotation of gene product function from high-throughput studies using the Gene Ontology. <i>Database: the Journal of Biological Databases and Curation</i> , 2019, 2019, .	1.4	21
23	JaponicusDB: rapid deployment of a model organism database for an emerging model species. <i>Genetics</i> , 2022, 220, .	1.2	21
24	The Protein Feature Ontology: a tool for the unification of protein feature annotations. <i>Bioinformatics</i> , 2008, 24, 2767-2772.	1.8	19
25	Community curation in PomBase: enabling fission yeast experts to provide detailed, standardized, sharable annotation from research publications. <i>Database: the Journal of Biological Databases and Curation</i> , 2020, 2020, .	1.4	19
26	Representing Kidney Development Using the Gene Ontology. <i>PLoS ONE</i> , 2014, 9, e99864.	1.1	17
27	Expanding yeast knowledge online. , 1998, 14, 1453-1469.		11
28	Developing an Ontology. <i>Methods in Molecular Biology</i> , 2008, 452, 111-124.	0.4	9
29	Term Matrix: a novel Gene Ontology annotation quality control system based on ontology term co-annotation patterns. <i>Open Biology</i> , 2020, 10, 200149.	1.5	7
30	Standards and Ontologies for Functional Genomics 2. <i>Comparative and Functional Genomics</i> , 2004, 5, 618-622.	2.0	6
31	Standards and Ontologies for Functional Genomics: Towards Unified Ontologies for Biology and Biomedicine. <i>Comparative and Functional Genomics</i> , 2003, 4, 116-120.	2.0	4