

Antonia Lock

List of Publications by Citations

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Version: 2024-04-29

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

15
papers

2,182
citations

12
h-index

19
g-index

19
ext. papers

5,699
ext. citations

9.9
avg, IF

4.1
L-index

#	Paper	IF	Citations
15	UniProt: the universal protein knowledgebase in 2021. <i>Nucleic Acids Research</i> , 2021 , 49, D480-D489	20.1	1073
14	The Gene Ontology resource: enriching a GOld mine. <i>Nucleic Acids Research</i> , 2021 , 49, D325-D334	20.1	494
13	PomBase: a comprehensive online resource for fission yeast. <i>Nucleic Acids Research</i> , 2012 , 40, D695-9	20.1	221
12	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	20.1	89
11	PomBase 2015: updates to the fission yeast database. <i>Nucleic Acids Research</i> , 2015 , 43, D656-61	20.1	74
10	A method for increasing expressivity of Gene Ontology annotations using a compositional approach. <i>BMC Bioinformatics</i> , 2014 , 15, 155	3.6	60
9	FYPO: the fission yeast phenotype ontology. <i>Bioinformatics</i> , 2013 , 29, 1671-8	7.2	35
8	Hidden in plain sight: what remains to be discovered in the eukaryotic proteome?. <i>Open Biology</i> , 2019 , 9, 180241	7	30
7	Canto: an online tool for community literature curation. <i>Bioinformatics</i> , 2014 , 30, 1791-2	7.2	30
6	Model organism databases: essential resources that need the support of both funders and users. <i>BMC Biology</i> , 2016 , 14, 49	7.3	30
5	PomBase: The Scientific Resource for Fission Yeast. <i>Methods in Molecular Biology</i> , 2018 , 1757, 49-68	1.4	15
4	One motif to bind them: A small-XXX-small motif affects transmembrane domain 1 oligomerization, function, localization, and cross-talk between two yeast GPCRs. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2014 , 1838, 3036-51	3.8	14
3	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,	5	10
2	Term Matrix: a novel Gene Ontology annotation quality control system based on ontology term co-annotation patterns. <i>Open Biology</i> , 2020 , 10, 200149	7	2
1	Term Matrix: A novel Gene Ontology annotation quality control system based on ontology term co-annotation patterns		1