

# Antonia Lock

## List of Publications by Year in Descending Order

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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	The Gene Ontology resource: enriching a GOld mine. <i>Nucleic Acids Research</i> , <b>2021</b> , 49, D325-D334	20.1	494
14	UniProt: the universal protein knowledgebase in 2021. <i>Nucleic Acids Research</i> , <b>2021</b> , 49, D480-D489	20.1	1073
13	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> , <b>2020</b> , 2020,	5	10
12	Term Matrix: a novel Gene Ontology annotation quality control system based on ontology term co-annotation patterns. <i>Open Biology</i> , <b>2020</b> , 10, 200149	7	2
11	PomBase 2018: user-driven reimplementaion of the fission yeast database provides rapid and intuitive access to diverse, interconnected information. <i>Nucleic Acids Research</i> , <b>2019</b> , 47, D821-D827	20.1	89
10	Hidden in plain sight: what remains to be discovered in the eukaryotic proteome?. <i>Open Biology</i> , <b>2019</b> , 9, 180241	7	30
9	PomBase: The Scientific Resource for Fission Yeast. <i>Methods in Molecular Biology</i> , <b>2018</b> , 1757, 49-68	1.4	15
8	Model organism databases: essential resources that need the support of both funders and users. <i>BMC Biology</i> , <b>2016</b> , 14, 49	7.3	30
7	PomBase 2015: updates to the fission yeast database. <i>Nucleic Acids Research</i> , <b>2015</b> , 43, D656-61	20.1	74
6	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> , <b>2014</b> , 1838, 3036-51	3.8	14
5	A method for increasing expressivity of Gene Ontology annotations using a compositional approach. <i>BMC Bioinformatics</i> , <b>2014</b> , 15, 155	3.6	60
4	Canto: an online tool for community literature curation. <i>Bioinformatics</i> , <b>2014</b> , 30, 1791-2	7.2	30
3	FYPO: the fission yeast phenotype ontology. <i>Bioinformatics</i> , <b>2013</b> , 29, 1671-8	7.2	35
2	PomBase: a comprehensive online resource for fission yeast. <i>Nucleic Acids Research</i> , <b>2012</b> , 40, D695-9	20.1	221
1	Term Matrix: A novel Gene Ontology annotation quality control system based on ontology term co-annotation patterns		1