

Sarah Hunter

List of Publications by Citations

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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.

16 papers	7,472 citations	11 h-index	16 g-index
16 ext. papers	9,983 ext. citations	12.6 avg, IF	4.65 L-index

#	Paper	IF	Citations
16	InterProScan 5: genome-scale protein function classification. <i>Bioinformatics</i> , 2014 , 30, 1236-40	7.2	3575
15	InterPro: the integrative protein signature database. <i>Nucleic Acids Research</i> , 2009 , 37, D211-5	20.1	1379
14	The InterPro protein families database: the classification resource after 15 years. <i>Nucleic Acids Research</i> , 2015 , 43, D213-21	20.1	954
13	InterPro in 2011: new developments in the family and domain prediction database. <i>Nucleic Acids Research</i> , 2012 , 40, D306-12	20.1	844
12	New developments in the InterPro database. <i>Nucleic Acids Research</i> , 2007 , 35, D224-8	20.1	397
11	EBI metagenomics--a new resource for the analysis and archiving of metagenomic data. <i>Nucleic Acids Research</i> , 2014 , 42, D600-6	20.1	104
10	Manual GO annotation of predictive protein signatures: the InterPro approach to GO curation. <i>Database: the Journal of Biological Databases and Curation</i> , 2012 , 2012, bar068	5	70
9	Genome3D: a UK collaborative project to annotate genomic sequences with predicted 3D structures based on SCOP and CATH domains. <i>Nucleic Acids Research</i> , 2013 , 41, D499-507	20.1	48
8	InterPro protein classification. <i>Methods in Molecular Biology</i> , 2011 , 694, 37-47	1.4	46
7	Meeting Report: BioSharing at ISMB 2010. <i>Standards in Genomic Sciences</i> , 2010 , 3, 254-8		18
6	Metagenomic analysis: the challenge of the data bonanza. <i>Briefings in Bioinformatics</i> , 2012 , 13, 743-6	13.4	16
5	The InterPro BioMart: federated query and web service access to the InterPro Resource. <i>Database: the Journal of Biological Databases and Curation</i> , 2011 , 2011, bar033	5	9
4	Meeting Report from the Genomic Standards Consortium (GSC) Workshop 10. <i>Standards in Genomic Sciences</i> , 2010 , 3, 225-31		7
3	Meeting Report: Metagenomics, Metadata and MetaAnalysis (M3) at ISMB 2010. <i>Standards in Genomic Sciences</i> , 2010 , 3, 232-4		3
2	MyDas, an extensible Java DAS server. <i>PLoS ONE</i> , 2012 , 7, e44180	3.7	2
1	Design, Implementation and Updating of Knowledge Bases 2010 , 87-105		