

Sarah Hunter

List of Publications by Year in descending order

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15
papers

11,254
citations

840776

11
h-index

996975

15
g-index

16
all docs

16
docs citations

16
times ranked

21108
citing authors

#	ARTICLE	IF	CITATIONS
1	InterProScan 5: genome-scale protein function classification. <i>Bioinformatics</i> , 2014, 30, 1236-1240.	4.1	6,553
2	InterPro: the integrative protein signature database. <i>Nucleic Acids Research</i> , 2009, 37, D211-D215.	14.5	1,712
3	The InterPro protein families database: the classification resource after 15 years. <i>Nucleic Acids Research</i> , 2015, 43, D213-D221.	14.5	1,205
4	InterPro in 2011: new developments in the family and domain prediction database. <i>Nucleic Acids Research</i> , 2012, 40, D306-D312.	14.5	921
5	New developments in the InterPro database. <i>Nucleic Acids Research</i> , 2007, 35, D224-D228.	14.5	444
6	EBI metagenomics—a new resource for the analysis and archiving of metagenomic data. <i>Nucleic Acids Research</i> , 2014, 42, D600-D606.	14.5	127
7	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-bar068.	3.0	108
8	InterPro Protein Classification. <i>Methods in Molecular Biology</i> , 2011, 694, 37-47.	0.9	63
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> , 2012, 41, D499-D507.	14.5	53
10	Metagenomic analysis: the challenge of the data bonanza. <i>Briefings in Bioinformatics</i> , 2012, 13, 743-746.	6.5	23
11	Meeting Report: BioSharing at ISMB 2010. <i>Standards in Genomic Sciences</i> , 2010, 3, 254-258.	1.5	19
12	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-bar033.	3.0	12
13	Meeting Report from the Genomic Standards Consortium (GSC) Workshop 10. <i>Standards in Genomic Sciences</i> , 2010, 3, 225-231.	1.5	8
14	Meeting Report: Metagenomics, Metadata and MetaAnalysis (M3) at ISMB 2010. <i>Standards in Genomic Sciences</i> , 2010, 3, 232-234.	1.5	4
15	MyDas, an Extensible Java DAS Server. <i>PLoS ONE</i> , 2012, 7, e44180.	2.5	2