

Emilie Pasche

List of Publications by Year in descending order

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Version: 2024-02-01

24
papers

321
citations

1306789

7
h-index

887659

17
g-index

27
all docs

27
docs citations

27
times ranked

701
citing authors

#	ARTICLE	IF	CITATIONS
1	Variomes: a high recall search engine to support the curation of genomic variants. <i>Bioinformatics</i> , 2022, 38, 2595-2601.	1.8	4
2	DisProt: intrinsic protein disorder annotation in 2020. <i>Nucleic Acids Research</i> , 2020, 48, D269-D276.	6.5	141
3	UPCLASS: a deep learning-based classifier for UniProtKB entry publications. <i>Database: the Journal of Biological Databases and Curation</i> , 2020, 2020, .	1.4	7
4	SIB Literature Services: RESTful customizable search engines in biomedical literature, enriched with automatically mapped biomedical concepts. <i>Nucleic Acids Research</i> , 2020, 48, W12-W16.	6.5	9
5	Accelerating annotation of articles via automated approaches: evaluation of the neXtA5 curation-support tool by neXtProt. <i>Database: the Journal of Biological Databases and Curation</i> , 2018, 2018, .	1.4	6
6	Triage by ranking to support the curation of protein interactions. <i>Database: the Journal of Biological Databases and Curation</i> , 2017, 2017, .	1.4	10
7	neXtA5: accelerating annotation of articles via automated approaches in neXtProt. <i>Database: the Journal of Biological Databases and Curation</i> , 2016, 2016, bau098.	1.4	10
8	Deep Question Answering for protein annotation. <i>Database: the Journal of Biological Databases and Curation</i> , 2015, 2015, bav081.	1.4	17
9	Development and tuning of an original search engine for patent libraries in medicinal chemistry. <i>BMC Bioinformatics</i> , 2014, 15, S15.	1.2	6
10	Closing the loop: from paper to protein annotation using supervised Gene Ontology classification. <i>Database: the Journal of Biological Databases and Curation</i> , 2014, 2014, bau088-bau088.	1.4	7
11	Managing the data deluge: data-driven GO category assignment improves while complexity of functional annotation increases. <i>Database: the Journal of Biological Databases and Curation</i> , 2013, 2013, bat041.	1.4	23
12	Assisted Knowledge Discovery for the Maintenance of Clinical Guidelines. <i>PLoS ONE</i> , 2013, 8, e62874.	1.1	5
13	Use of controlled vocabularies to improve biomedical information retrieval tasks. <i>Studies in Health Technology and Informatics</i> , 2013, 192, 1068.	0.2	1
14	Using binary classification to prioritize and curate articles for the Comparative Toxicogenomics Database. <i>Database: the Journal of Biological Databases and Curation</i> , 2012, 2012, bas050-bas050.	1.4	7
15	Answering Gene Ontology terms to proteomics questions by supervised macro reading in Medline. <i>EMBnet Journal</i> , 2012, 18, 29.	0.2	6
16	Development of a text search engine for medicinal chemistry patents. <i>EMBnet Journal</i> , 2012, 18, 44.	0.2	1
17	Building a Transnational Biosurveillance Network Using Semantic Web Technologies: Requirements, Design, and Preliminary Evaluation. <i>Journal of Medical Internet Research</i> , 2012, 14, e73.	2.1	29
18	Interoperability driven integration of biomedical data sources. <i>Studies in Health Technology and Informatics</i> , 2011, 169, 185-9.	0.2	10

#	ARTICLE	IF	CITATIONS
19	Using multimodal mining to drive clinical guidelines development. Studies in Health Technology and Informatics, 2011, 169, 477-81.	0.2	1
20	Simple Pre and Post Processing Strategies for Patent Searching in CLEF Intellectual Property Track 2009. Lecture Notes in Computer Science, 2010, , 444-451.	1.0	4
21	QA-driven guidelines generation for bacteriotherapy. AMIA ... Annual Symposium proceedings, 2009, 2009, 509-13.	0.2	2
22	DebugIT: building a European distributed clinical data mining network to foster the fight against microbial diseases. Studies in Health Technology and Informatics, 2009, 148, 50-9.	0.2	4
23	Biomedical data management: a proposal framework. Studies in Health Technology and Informatics, 2009, 150, 175-9.	0.2	6
24	Automatic medical knowledge acquisition using question-answering. Studies in Health Technology and Informatics, 2009, 150, 569-73.	0.2	4