

# Emilie Pasche

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6758824/publications.pdf>

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  | DisProt: intrinsic protein disorder annotation in 2020. <i>Nucleic Acids Research</i> , 2020, 48, D269-D276.   | 6.5 | 141       |
| 2  | 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        |
| 3  | 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        |
| 4  | Deep Question Answering for protein annotation. <i>Database: the Journal of Biological Databases and Curation</i> , 2015, 2015, bav081.  | 1.4 | 17        |
| 5  | neXtA5: accelerating annotation of articles via automated approaches in neXtProt. <i>Database: the Journal of Biological Databases and Curation</i> , 2016, 2016, baw098.  | 1.4 | 10        |
| 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  | Interoperability driven integration of biomedical data sources. <i>Studies in Health Technology and Informatics</i> , 2011, 169, 185-9.  | 0.2 | 10        |
| 8  | 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         |
| 9  | 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         |
| 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 | 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         |
| 12 | Development and tuning of an original search engine for patent libraries in medicinal chemistry. <i>BMC Bioinformatics</i> , 2014, 15, S15.  | 1.2 | 6         |
| 13 | 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         |
| 14 | Answering Gene Ontology terms to proteomics questions by supervised macro reading in Medline. <i>EMBnet Journal</i> , 2012, 18, 29.  | 0.2 | 6         |
| 15 | Biomedical data management: a proposal framework. <i>Studies in Health Technology and Informatics</i> , 2009, 150, 175-9.  | 0.2 | 6         |
| 16 | Assisted Knowledge Discovery for the Maintenance of Clinical Guidelines. <i>PLoS ONE</i> , 2013, 8, e62874.  | 1.1 | 5         |
| 17 | Simple Pre and Post Processing Strategies for Patent Searching in CLEF Intellectual Property Track 2009. <i>Lecture Notes in Computer Science</i> , 2010, , 444-451.   | 1.0 | 4         |
| 18 | DebugIT: building a European distributed clinical data mining network to foster the fight against microbial diseases. <i>Studies in Health Technology and Informatics</i> , 2009, 148, 50-9.                       | 0.2 | 4         |

| #  | ARTICLE  | IF  | CITATIONS |
|----|--|-----|-----------|
| 19 | Automatic medical knowledge acquisition using question-answering. <i>Studies in Health Technology and Informatics</i> , 2009, 150, 569-73.               | 0.2 | 4         |
| 20 | Variomes: a high recall search engine to support the curation of genomic variants. <i>Bioinformatics</i> , 2022, 38, 2595-2601.                          | 1.8 | 4         |
| 21 | QA-driven guidelines generation for bacteriotherapy. <i>AMIA ... Annual Symposium proceedings</i> , 2009, 2009, 509-13.                                  | 0.2 | 2         |
| 22 | Development of a text search engine for medicinal chemistry patents. <i>EMBnet Journal</i> , 2012, 18, 44.   | 0.2 | 1         |
| 23 | Using multimodal mining to drive clinical guidelines development. <i>Studies in Health Technology and Informatics</i> , 2011, 169, 477-81.               | 0.2 | 1         |
| 24 | Use of controlled vocabularies to improve biomedical information retrieval tasks. <i>Studies in Health Technology and Informatics</i> , 2013, 192, 1068. | 0.2 | 1         |