

# Robert Stevens

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5285939/publications.pdf>

Version: 2024-02-01

34  
papers

1,160  
citations

623574

14  
h-index

454834

30  
g-index

35  
all docs

35  
docs citations

35  
times ranked

1472  
citing authors

#	ARTICLE	IF	CITATIONS
1	NERO: a biomedical named-entity (recognition) ontology with a large, annotated corpus reveals meaningful associations through text embedding. <i>Npj Systems Biology and Applications</i> , 2021, 7, 38.	1.4	3
2	Semantic Deep Learning: Prior Knowledge and a Type of Four-Term Embedding Analogy to Acquire Treatments for Well-Known Diseases. <i>JMIR Medical Informatics</i> , 2020, 8, e16948.	1.3	0
3	Exploring semantic deep learning for building reliable and reusable one health knowledge from PubMed systematic reviews and veterinary clinical notes. <i>Journal of Biomedical Semantics</i> , 2019, 10, 22.	0.9	6
4	Measuring expert performance at manually classifying domain entities under upper ontology classes. <i>Web Semantics</i> , 2019, 57, 100469.	2.2	7
5	Unveiling antimicrobial peptideâ€“generating human proteases using PROTEASIX. <i>Journal of Proteomics</i> , 2018, 171, 53-62.	1.2	11
6	Inference Inspector: Improving the verification of ontology authoring actions. <i>Web Semantics</i> , 2018, 49, 1-15.	2.2	12
7	Deep learning meets ontologies: experiments to anchor the cardiovascular disease ontology in the biomedical literature. <i>Journal of Biomedical Semantics</i> , 2018, 9, 13.	0.9	28
8	MIRO: guidelines for minimum information for the reporting of an ontology. <i>Journal of Biomedical Semantics</i> , 2018, 9, 6.	0.9	55
9	Prediction of Proteases Involved in Peptide Generation. <i>Methods in Molecular Biology</i> , 2017, 1574, 205-213.	0.4	2
10	A Case Study on Sepsis Using PubMed and Deep Learning for Ontology Learning. <i>Studies in Health Technology and Informatics</i> , 2017, 235, 516-520.	0.2	3
11	The BioHub Knowledge Base: Ontology and Repository for Sustainable Biosourcing. <i>Journal of Biomedical Semantics</i> , 2016, 7, 30.	0.9	2
12	The Proteasix Ontology. <i>Journal of Biomedical Semantics</i> , 2016, 7, 33.	0.9	8
13	Supporting the analysis of ontology evolution processes through the combination of static and dynamic scaling functions in OQuARE. <i>Journal of Biomedical Semantics</i> , 2016, 7, 63.	0.9	8
14	Omics databases on kidney disease: where they can be found and how to benefit from them. <i>CKJ: Clinical Kidney Journal</i> , 2016, 9, 343-352.	1.4	33
15	Ten Simple Rules for Selecting a Bio-ontology. <i>PLoS Computational Biology</i> , 2016, 12, e1004743.	1.5	29
16	Evaluating the Emotion Ontology through use in the self-reporting of emotional responses at an academic conference. <i>Journal of Biomedical Semantics</i> , 2014, 5, 38.	0.9	14
17	The Software Ontology (SWO): a resource for reproducibility in biomedical data analysis, curation and digital preservation. <i>Journal of Biomedical Semantics</i> , 2014, 5, 25.	0.9	56
18	The Quality of Methods Reporting in Parasitology Experiments. <i>PLoS ONE</i> , 2014, 9, e101131.	1.1	12

#	ARTICLE	IF	CITATIONS
19	Stealthy annotation of experimental biology by spreadsheets. <i>Concurrency Computation Practice and Experience</i> , 2013, 25, 467-480.	1.4	3
20	Three Steps to Heaven: Semantic Publishing in a Real World Workflow. <i>Future Internet</i> , 2012, 4, 1004-1015.	2.4	1
21	Engineering use cases for modular development of ontologies in OWL. <i>Applied Ontology</i> , 2012, 7, 113-132.	1.0	18
22	Using semantic web technologies to manage complexity and change in biomedical data. , 2011, 2011, 3708-11.		2
23	Building Workflows that Traverse the Bioinformatics Data Landscape. , 2009, , 141-163.		0
24	BioCatalogue: A Curated Web Service Registry For The Life Science Community. <i>Nature Precedings</i> , 2009, , .	0.1	13
25	Mining Taverna's semantic web of provenance. <i>Concurrency Computation Practice and Experience</i> , 2008, 20, 463-472.	1.4	85
26	Process of Building a Vocabulary for the Infection Domain. , 2008, , .		9
27	Using provenance to manage knowledge of In Silico experiments. <i>Briefings in Bioinformatics</i> , 2007, 8, 183-194.	3.2	31
28	Taverna: lessons in creating a workflow environment for the life sciences. <i>Concurrency Computation Practice and Experience</i> , 2006, 18, 1067-1100.	1.4	485
29	Augmenting the mobility of profoundly blind Web travellers. <i>New Review of Hypermedia and Multimedia</i> , 2005, 11, 103-128.	0.9	11
30	Building Ontologies in DAML + OIL. <i>Comparative and Functional Genomics</i> , 2003, 4, 133-141.	2.0	17
31	A SUITE OF DAML+OIL ONTOLOGIES TO DESCRIBE BIOINFORMATICS WEB SERVICES AND DATA. <i>International Journal of Cooperative Information Systems</i> , 2003, 12, 197-224.	0.6	126
32	OILing the way to machine understandable bioinformatics resources. <i>IEEE Transactions on Information Technology in Biomedicine</i> , 2002, 6, 129-134.	3.6	21
33	Building a bioinformatics ontology using OIL. <i>IEEE Transactions on Information Technology in Biomedicine</i> , 2002, 6, 135-141.	3.6	44
34	Ontology Based Document Enrichment in Bioinformatics. <i>Comparative and Functional Genomics</i> , 2002, 3, 42-46.	2.0	5