

OrganismTagger: Detection, normalization, and grounding of organism entities in biomedical documents

Submitted by [rene](#) [1] on Sun, 2011-08-14 18:00

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Title	OrganismTagger: Detection, normalization, and grounding of organism entities in biomedical documents
Publication Type	Journal Article
Year of Publication	2011
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Refereed Designation	Refereed
Journal	Bioinformatics
Volume	27
Issue	19
Pagination	2721--2729
Date Published	August 9, 2011
ISSN	1460-2059 (online) 1367-4803 (print)
Abstract	<p>Motivation: Semantic tagging of organism mentions in full-text articles is an important part of literature mining and semantic enrichment solutions. Tagged organism mentions also play a pivotal role in disambiguating other entities in a text, such as proteins. A high-precision organism tagging system must be able to detect the numerous forms of organism mentions, including common names as well as the traditional taxonomic groups: genus, species, and strains. In addition, such a system must resolve abbreviations and acronyms, assign the scientific name, and if possible link the detected mention to the NCBI Taxonomy database for further semantic queries and literature navigation.</p> <p>Results: We present the OrganismTagger, a hybrid rule-based/machine learning system to extract organism mentions from the literature. It includes tools for automatically generating lexical and ontological resources from a copy of the NCBI Taxonomy database, thereby facilitating system updates by end-users. Its novel ontology-based resources can also be reused in other semantic mining and linked data tasks. Each detected organism mention is normalized to a canonical name through the resolution of acronyms and abbreviations and subsequently grounded with an NCBI Taxonomy database ID. In particular, our system combines a novel machine-learning approach with rule-based and lexical methods for detecting strain mentions in documents. On our manually annotated OT corpus, the OrganismTagger achieves a precision of 95%, a recall of 94% and a grounding accuracy of 97.5%. On the manually annotated corpus of Linnaeus-100, the results show a precision of 99%, recall of 97% and grounding accuracy of 97.4%.</p>

Availability: The OrganismTagger, including supporting tools, resources, training data and manual annotations, as well as end-user and developer documentation, is freely available under an open source license at

<http://www.semanticsoftware.info/organism-tagger> [9].

URL	http://bioinformatics.oxfordjournals.org/cgi/content/abstract/btr452?jkey=9uBtAIDN8y2nmz6&keytype=ref [10]
DOI	10.1093/bioinformatics/btr452 [11]
Impact Factor	5.468 (2012)
History	Received on March 7, 2011; revised on July 14, 2011; accepted on July 31, 2011.
Attachment	Size
Bioinformatics-2011-Naderi-2721-9.pdf [12]	424.64 KB



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