

## RESEARCH

## A sample article title

Jane E Doe<sup>1\*†</sup> and John RS Smith<sup>1,2</sup>**Abstract****First part title:** Text for this section.**Second part title:** Text for this section.**Keywords:** sample; article; author**Content**

Text and results for this section, as per the individual journal's instructions for authors.

**Background**

Incessant influx of new data on chemical entities and their properties steadily increases our capabilities for information retrieval in biomedical domain. However, ever-growing rates at which new information is being produced make it infeasible for any specialist to analyze manually even if confined within a specific subject. On the other hand the prevalent formats for such publication (paper, theses or a patent) is not structured in a way that is suitable for automatic processing[1]. Various databases containing diverse chemical information, such as PubChem[2], ChemSpider[3], CAS Registry[4], etc., partially address this problem by accumulating information from different sources and storing it in an organized, easy-to-process manner. But the maintenance and curation of this kind of databases still requires substantial manual labour from the experts in the field. Moreover, a standardized entry to a database usually omits many of the nuances transmitted by a natural language. For this reason, in recent years special interest is given to the instruments for automatic information extraction from text[5].

Named entities recognition (NER) is an essential step in the process of extracting chemical information from text. The problem of NER is not unique to chemical entities, but there are several features characteristic to the given subject. A lack of universal naming convention, the existence of multiple synonyms for a single entity (e.g. EXAMPLE), high sensitivity to

spelling errors and a hard-to-parse syntax with multiple whitespaces, punctuation marks and numbers inside single entity name are some of the most obvious examples[6, 7]. Significance of text mining and information extraction in biomedical domain is outlined by such initiatives as BioCreative challenge[8] and chemical named entities recognition (CNER) in particular by CHEMDNER task[9].

Historically, first approaches used for CNER were dictionary-based and rule-based. Dictionary-based algorithms rely on string matching to a domain-specific collection of terms. They are highly dependent on the quality of this term collection, which can get outdated in time and hence requires constant expansion and curation[10, 11]. Rule-based algorithms make use of manually created and updated rules based on morphological and contextual information. These rules allow users to describe potentially endless set of names if they follow an orderly convention such as IUPAC, but do not port as well on trivial names and like dictionaries require maintenance by the experts[12].

**Competing interests**

The authors declare that they have no competing interests.

**Author's contributions**

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**Acknowledgements**

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**Author details**

<sup>1</sup>Department of Zoology, Cambridge, Waterloo Road, London, UK.

<sup>2</sup>Marine Ecology Department, Institute of Marine Sciences Kiel, Düsternbrooker Weg 20, 24105 Kiel, Germany.

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\*Correspondence: jane.e.doe@cambridge.co.uk

<sup>1</sup>Department of Zoology, Cambridge, Waterloo Road, London, UK

Full list of author information is available at the end of the article

<sup>†</sup>Equal contributor

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## Figures

**Figure 1 Sample figure title.** A short description of the figure content should go here.

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## Tables

**Table 1** Sample table title. This is where the description of the table should go.

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Additional file 1 — Sample additional file title

Additional file descriptions text (including details of how to view the file, if it is in a non-standard format or the file extension). This might refer to a multi-page table or a figure.

Additional file 2 — Sample additional file title

Additional file descriptions text.