

City Research Online

City, University of London Institutional Repository

Citation: Jimenez-Ruiz, E. (2023). LogMap Family Participation in the OAEI 2023. OM 2023 Ontology Matching, 3591, pp. 157-158. ISSN 1613-0073

This is the published version of the paper.

This version of the publication may differ from the final published version.

Permanent repository link: https://openaccess.city.ac.uk/id/eprint/32010/

Link to published version:

Copyright: City Research Online aims to make research outputs of City, University of London available to a wider audience. Copyright and Moral Rights remain with the author(s) and/or copyright holders. URLs from City Research Online may be freely distributed and linked to.

Reuse: Copies of full items can be used for personal research or study, educational, or not-for-profit purposes without prior permission or charge. Provided that the authors, title and full bibliographic details are credited, a hyperlink and/or URL is given for the original metadata page and the content is not changed in any way.

LogMap Family Participation in the OAEI 2023

Ernesto Jiménez-Ruiz

¹Department of Computer Science, City, University of London, UK ²Department of Informatics, University of Oslo, Oslo, Norway

Abstract

We present the participation of LogMap and its variants in the OAEI 2023 campaign. The LogMap project started in January 2011 with the objective of developing a scalable and logic-based ontology matching system.

1. Presentation of the system

LogMap [1, 2] is an ontology matching system that *(i)* can efficiently match semantically rich ontologies containing tens (and even hundreds) of thousands of classes, *(ii)* incorporates sophisticated reasoning and repair techniques to minimise the number of logical inconsistencies [3], and *(iii)* provides support for user intervention during the matching process [4]. LogMap ISWC 2011 paper [1] has recently been awarded the SWSA Ten-Year Award.¹

1.1. LogMap variants in the 2023 campaign

As in previous campaigns, in the OAEI 2023 we have participated with two additional variants:

- **LogMapLt** is a "lightweight" variant of LogMap, which essentially only applies (efficient) string matching techniques.
- **LogMapBio** includes an extension to use BioPortal [5] as a (dynamic) provider of mediating ontologies instead of relying on a few preselected ontologies [6].

1.2. Link to the system and parameters file

LogMap is open-source and released under the Apache-2.0 License.² LogMap components and source code are available from the LogMap's GitHub page: https://github.com/ernestojimenezruiz/logmap-matcher/.

OM 2023: The 18th International Workshop on Ontology Matching collocated with the 22nd International Semantic Web Conference ISWC-2023 November 7th, 2023, Athens, Greece

ernesto.jimenez-ruiz@city.ac.uk (E. Jiménez-Ruiz)

https://www.city.ac.uk/about/people/academics/ernesto-jimenez-ruiz (E. Jiménez-Ruiz)

D 0000-0002-9083-4599 (E. Jiménez-Ruiz)

^{© 023} Copyright for this paper by its authors. Use permitted under Creative Commons License Attribution 4.0 International (CC BY 4.0).

CEUR Workshop Proceedings (CEUR-WS.org)

¹http://swsa.semanticweb.org/content/swsa-ten-year-award

²http://www.apache.org/licenses/

LogMap distributions can be easily customized through a configuration file containing the matching parameters.

1.3. Results

Please refer to http://oaei.ontologymatching.org/2023/results/ for the results of the LogMap family in the OAEI 2023 campaign. It is worth mentioning that LogMap-Bio was one of the top systems in the new Bio-ML track [7, 8].

Acknowledgments

I would like to thank Bernardo Cuenca-Grau, Ian Horrocks, Alessandro Solimando, Jiaoyan Chen, Valerie Cross, Anton Morant, Yujiao Zhou, Weiguo Xia, Xi Chen, Yuan Gong, Shuo Zhang and Rob Upson, who have contributed to the LogMap project in the past.

References

- E. Jiménez-Ruiz, B. Cuenca Grau, LogMap: Logic-based and Scalable Ontology Matching, in: Int'l Sem. Web Conf. (ISWC), 2011, pp. 273–288.
- [2] E. Jiménez-Ruiz, B. Cuenca Grau, Y. Zhou, I. Horrocks, Large-scale interactive ontology matching: Algorithms and implementation, in: Europ. Conf. on Artif. Intell. (ECAI), 2012.
- [3] A. Solimando, E. Jiménez-Ruiz, G. Guerrini, Minimizing conservativity violations in ontology alignments: Algorithms and evaluation, Knowledge and Information Systems (2016).
- [4] H. Li, Z. Dragisic, D. Faria, V. Ivanova, E. Jiménez-Ruiz, P. Lambrix, C. Pesquita, User validation in ontology alignment: functional assessment and impact, Knowl. Eng. Rev. 34 (2019) e15. URL: https://doi.org/10.1017/S0269888919000080. doi:10.1017/S0269888919000080.
- [5] N. Fridman Noy, N. H. Shah, P. L. Whetzel, B. Dai, et al., BioPortal: ontologies and integrated data resources at the click of a mouse, Nucleic Acids Research 37 (2009) 170–173.
- [6] X. Chen, W. Xia, E. Jiménez-Ruiz, V. Cross, Extending an ontology alignment system with bioportal: a preliminary analysis, in: Poster at Int'l Sem. Web Conf. (ISWC), 2014.
- Y. He, J. Chen, H. Dong, E. Jiménez-Ruiz, A. Hadian, I. Horrocks, Machine Learning-Friendly Biomedical Datasets for Equivalence and Subsumption Ontology Matching, in: The Semantic Web - ISWC 2022 - 21st International Semantic Web Conference, Virtual Event, October 23-27, 2022, Proceedings, volume 13489 of *Lecture Notes in Computer Science*, Springer, 2022, pp. 575–591. URL: https://doi.org/10.1007/978-3-031-19433-7_33. doi:10. 1007/978-3-031-19433-7_33.
- [8] Y. He, J. Chen, H. Dong, E. Jiménez-Ruiz, A. Hadian, I. Horrocks, Bio-ML: Machine Learning-Friendly Biomedical Datasets for Equivalence and Subsumption Ontology Matching (Version OAEI Bio-ML 2023), https://doi.org/10.5281/zenodo.8193375, 2023. URL: https://doi.org/10. 5281/zenodo.8193375. doi:10.5281/ZENODO.8193375.