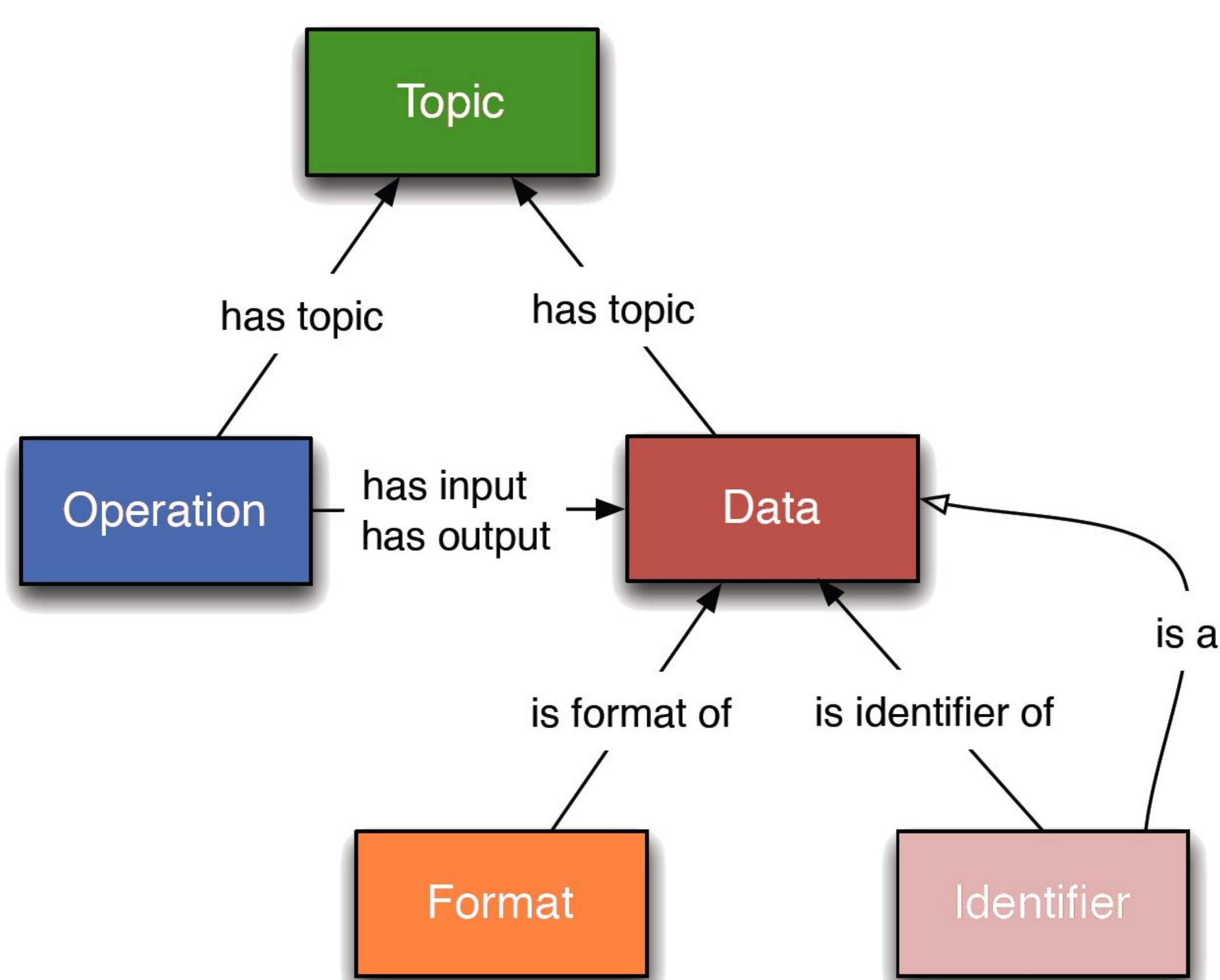


EDAM is an ontology of well-established, familiar concepts that are prevalent within bioinformatics, and life science data analysis in general. The scope of EDAM includes types of data (including identifiers), data formats, operations, and topics. EDAM has a relatively simple architecture, and comprises a set of concepts with terms, synonyms, definitions, relations, links, and some additional information (especially for formats).

Architecture



Overview



Open source and community-driven continuous development



Used in diverse applications and projects



... and many more

Available in various ontology browsers and download formats



Matúš Kalaš¹



Hervé Ménager²



Veit Schwämmle³



Jon Ison⁴

¹ Computational Biology Unit, Department of Informatics, University of Bergen, Norway
² Bioinformatics and Biostatistics Hub - C3BI, USR 3756 IP CNRS - Institut Pasteur, Paris, France
³ University of Southern Denmark, Ødense, Denmark
⁴ IT Service, Technical University of Denmark, Kongens Lyngby, Denmark