DataCite Services & Downstream Impact







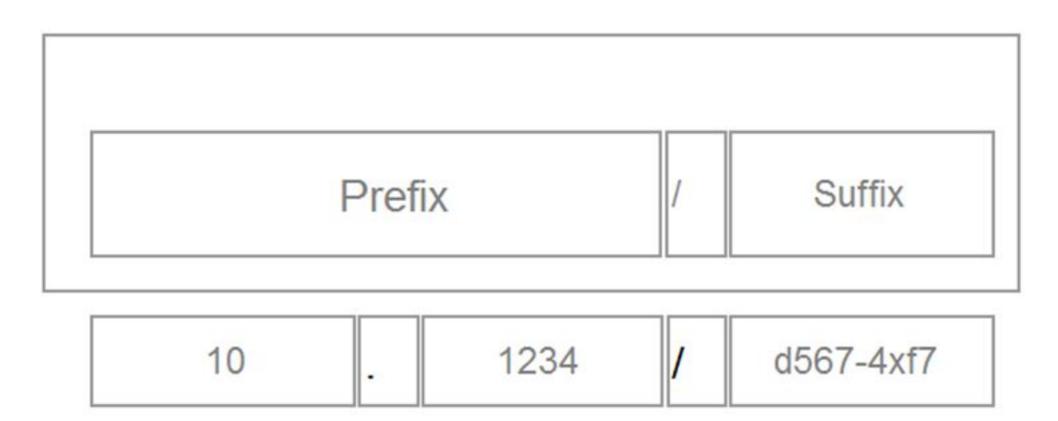
Britta Dreyer

https://orcid.org/0000-0002-0687-5460 Webinar, **Middle East DOI Consortium**, 11-May-2022

What is a DOI?



- Persistent Identifier and hyperlink
- Resolves via Google with any browser
- Specifies the resource not the location Landing page URL
- ☐ Has standard metadata attached DataCite metadata schema
- Making research FAIR (Findable, Accessible, Interoperable, Reusable)
- ☐ Standard DIN ISO 26324 (2012)





DOIs for Research Output

1. Take an article

2. Describe it

over all the classes that can occur in that position. Restik's model was proposed as a model of human learning of selectional preferences that made minimal representational assumptions; it showed how such preferences could be acquired from usage data and an existing conceptual hierarchy. However, his and later computational models (see Section 2) have properties that do not match with certain cognitive plausibility criteria for a child language acquisition model. All these models use the training data in "batch mode", and most of them use information theoretic measures that rely on total counts from a corpus. Therefore, it is not clear how the representation of selectional preferences could be updated in-



3. Assign a DOI 10.12345/h43t-z2hd

4. Reuse and reference

Citation

APA

Hosseini, B., Shahriari-Ahmadi, F., Hashemi, H., Marashi, M.-H., Mohseniazar, M., Farokhzad, A., & Sabokbari, M. (2011). Transient Expression of cor Gene in Papaver somniferum. BioImpacts; ISSN 2228-5660. https://doi.org/10.5681/BI.2011.033

5. Enjoy the benefits!

Findability Track citations Reusability Measure impact





Unique Persistent

DOIs for various Resource Types

- Research data
- Dissertations
- Journal articles
- Preprints
- Book chapters
- Images
- Videos
- Conference papers
- Samples
- □ Software...



Resource Type

☐ Dataset	12,804,731
☐ Text	10,019,200
☐ Image	3,180,638
Other	1,929,426
Physical Object	1,394,617
☐ Preprint	928,363
☐ Collection	797,640
Software	292,351
Audiovisual	266,883
☐ Journal Article	223,002
Report	81,473
Dissertation	62,762
Sound	40,279
☐ Interactive Resource	32,395
☐ Conference Paper	22,327
Book	15,013

Metadata Schema



Mandatory	Recommended	Optional
Identifier	Subject	Language
Creator (with ORCiD) *	Contributor	Alternate ID
Title	Date	Size
Publisher	Related identifier	Format
Publication year	Description	Version
Resource Type	GeoLocation	Rights

Current version 4.3 XML examples available

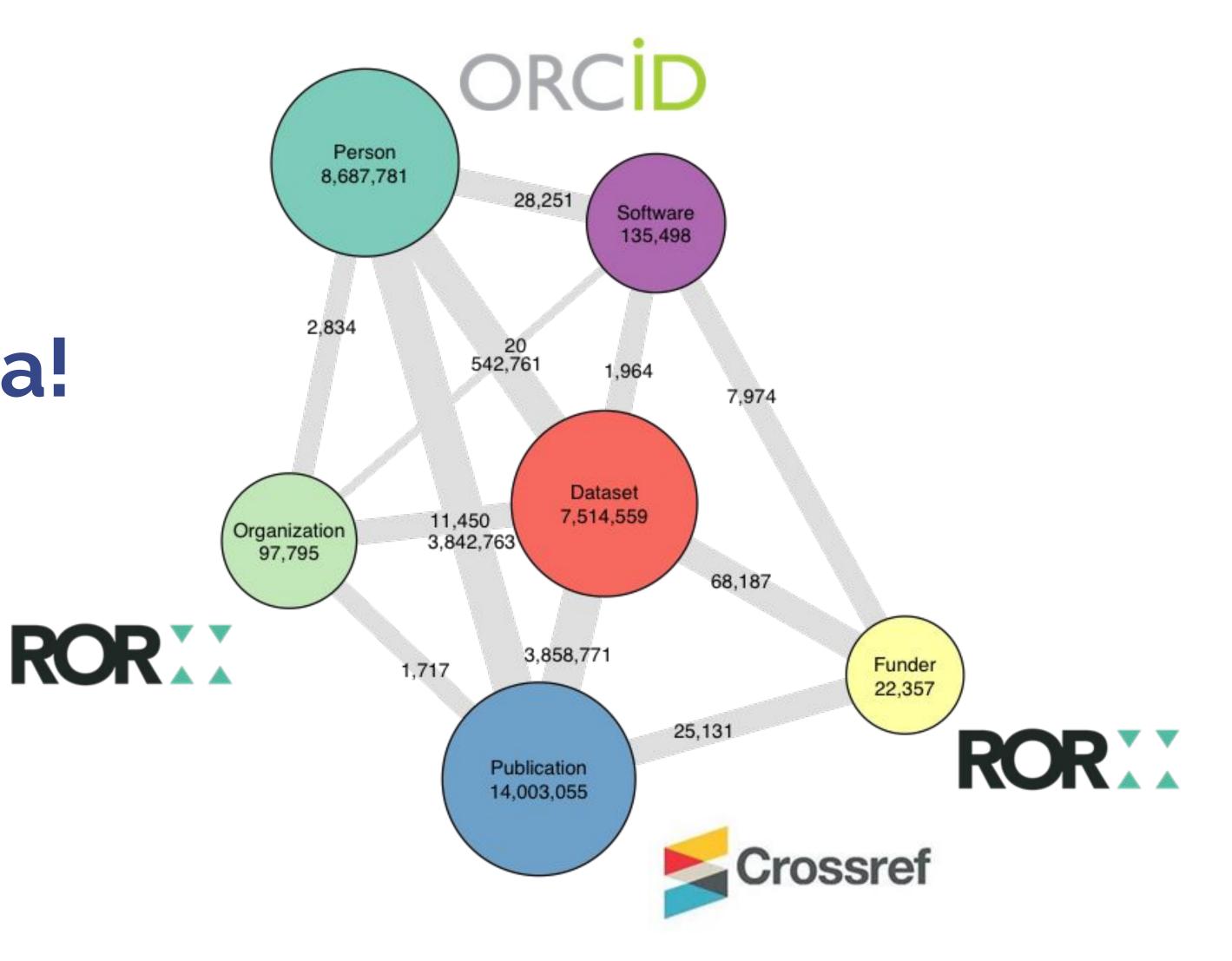
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RIS
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^{*} ORCID is optional

Open Science Research Graph



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PIDs to the
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Landing page



International Journal of Innovative Technology & Interdisciplinary Sciences (IJITIS) ISSN: 2613-7305, Peer-Reviewed, Quarterly

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□ DOI as full https URL

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Vol. 5 No. 2 (2022): International Journal of Innovative Technology and Interdisciplinary Sciences - Volume 5, Issue 2 / Articles

Weighted Trimean as a Regressor in the Estimate of Theil-Sen Regression

Necati Alp Erilli

Department of Econometrics, Sivas Cumhuriyet University, Sivas, Turkey

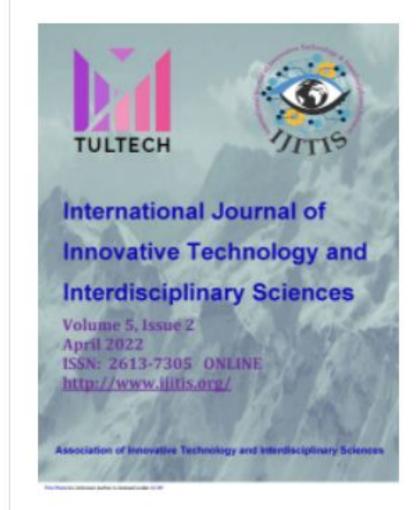
DOI: https://doi.org/10.15157/IJITIS.2022.5.2.892-906



Keywords: Theil-Sen Regression Analysis; weighted trimean; non-parametric regression analysis; trimean

Abstract

The most used method in nonparametric regression analysis is the Theil-Sen approach. With this method, all coefficient estimations are made with the median parameter as opposed to parametric methods. The most important criticism in computations with the median parameter is that the impact of extreme values does not participate in calculations. In this study, it was proposed to use the trimean parameter by weighting, which more effectively adds the effect of outliers to the average account in Theil-Sen regression analysis. In applications with 5 data sets, Theil-Sen calculations with weighted trimean were found to be more successful than calculations with the median parameter. Thus, in cases where the outliers are too high or directly affect the data, it can be said that the use of weighted trimean will yield more effective results.







Published

2022-03-16 - Updated on 2022-03-17

How to Cite



Necati Alp Erilli. (2022). Weighted Trimean as a Regressor in the Estimate of Theil-Sen Regression. International Journal of Innovative Technology and Interdisciplinary Sciences, 5(2), 892– 906. Make a Submission

Information

For Readers

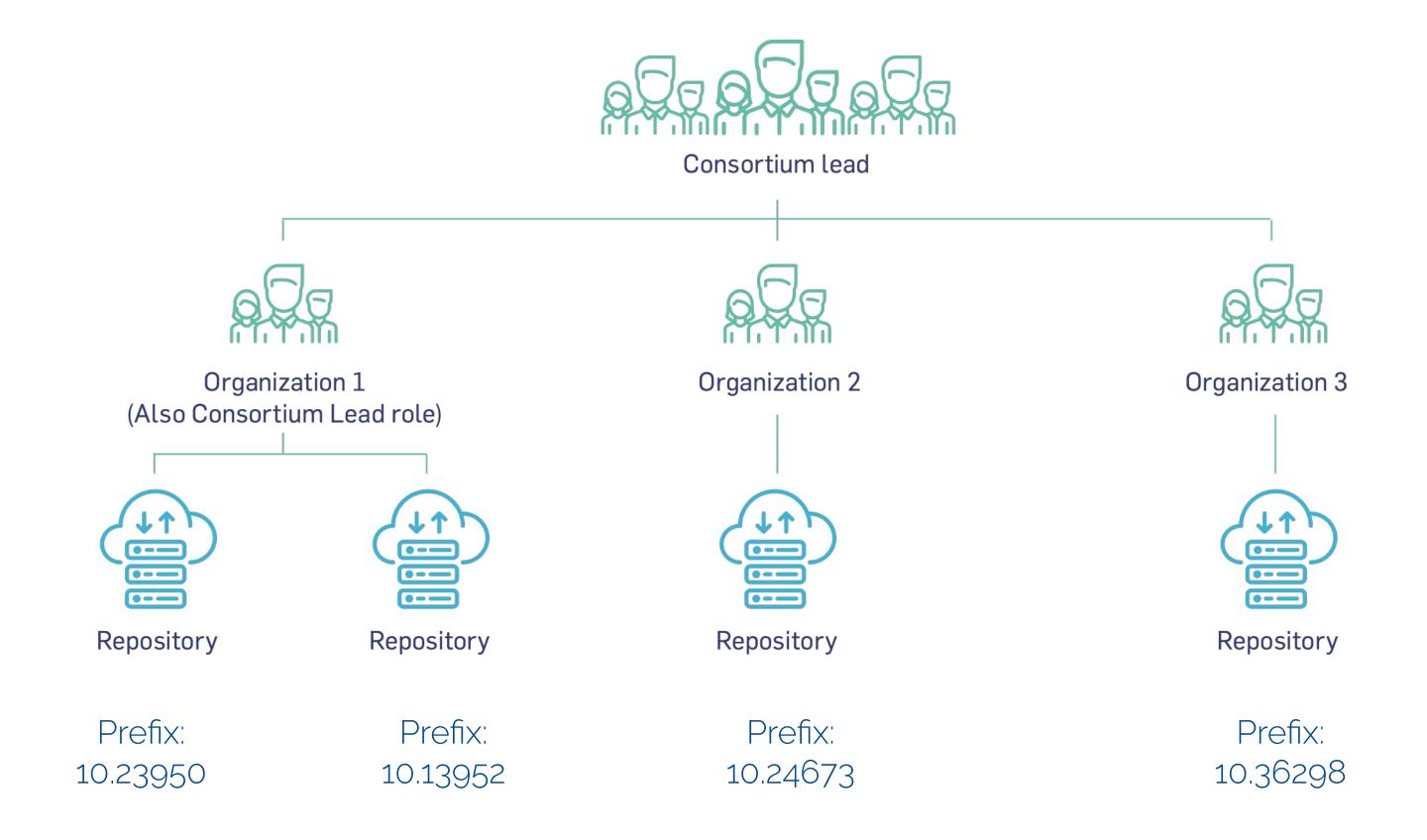
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DataCite Services

Consortium membership Middle East DOI Consortium





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Transfer

1,163 DOIs

Reset All

Type to search...

☐ Findable 1,163

Resource Type

State

☐ Text 1,087

Transient Expression of cor Gene in Papaver somniferum Text

Bahman Hosseini, Farajollah Shahriari-Ahmadi, Haleh Hashemi, Mohammad-Hasan Marashi, Mahdi Mohseniazar, Alireza Farokhzad, Masoume Sabokbari,

Journal Article published 2011 via BioImpacts; ISSN 2228-5660

Introduction: Papaver somniferum is the commercial source of morphine and codeine. The isolation of effective genes involved in the morphine biosynthesis of P. somniferum is very important in the production of specific metabolites achieved using metabolic engineering techniques. In this pathway, the key enzyme COR is involved in the conversion of codeinone to codeine and morphinone to

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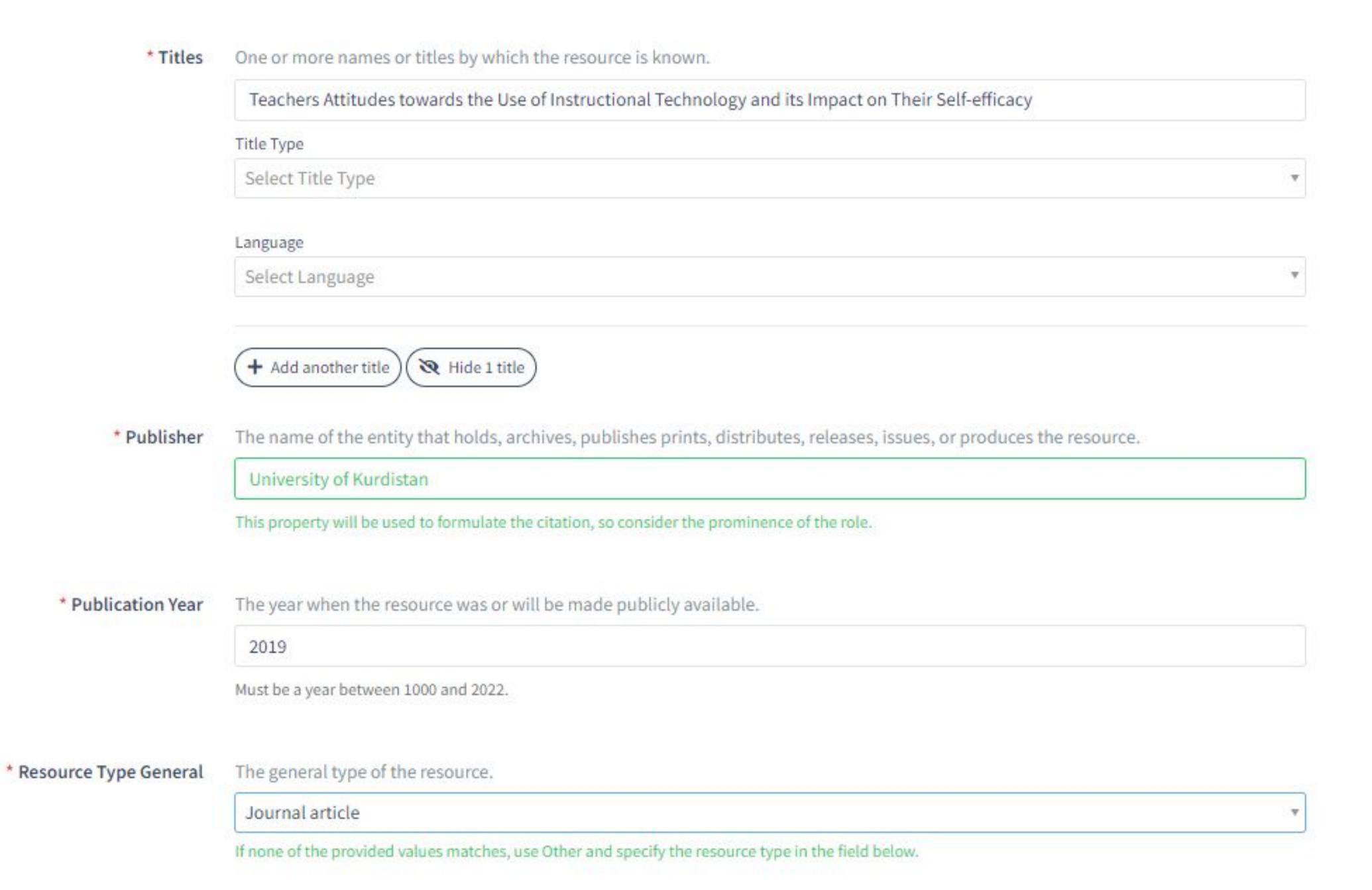
More information about DOI registration via form can be found on our Support Website. Required properties are marked with a red

Required Properties

asterix. The globally unique string that identifies the resource and can't be changed. 10.34785/j014.2022.579 The state determines whether a DOI is registered and findable. Once in Registered or Findable state, a DOI can't be set back to Draft state. More ... Draft only visible in Fabrica, DOI can be deleted O Registered registered with the DOI Resolver Findable registered with the DOI Resolver and indexed in DataCite Search The location of the landing page with more information about the resource. https://cls.uok.ac.ir/article_62222.html Should be a https URL - within the allowed domain(s) of your repository if domain restrictions are enabled in the repository settings. Http and ftp are also supported. The main researchers or organizations involved in producing the resource, in priority order. * Creators Name Identifier Name Identifier Uniquely identifies an individual or legal entity, according to various schemas, e.g. ORCID, ROR or ISNI. Use name identifier expressed as URL. The Given Name, Family Name and Name will automatically be filled out for ORCID and ROR identifiers. Add another name identifier Person Organization Unknown Given Name nick The personal or first name of the creator. Family Name حسيني The surname or last name of the creator.

سىنساجد، nick مسينى

* Name (from Given Name and Family Name)



Testing



- Test the complete workflow incl. DOI stages in the test environment and resolving to landingpages
- The same account credentials work for web interface and the API:
 - DOI Fabrica (https://doi.test.datacite.org)
 - MDS API (Test endpoint: https://mds.test.datacite.org)
 - JSON REST API (Test endpoint: https://api.test.datacite.org)
- · Assign or request a prefix to/for your repository account
- Testing guide: https://support.datacite.org/docs/testing-guide If you have any questions please contact support@datacite.org.

Support Services



Support Site: https://support.datacite.org

Support Desk: support@datacite.org



WEB INTERFACE - FABRICA

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- > Direct Members
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DataCite Metadata Schema 4.4

DataCite Metadata Schema 4.3

DataCite Metadata Schema 4.2

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DataCite DOI Display Guidelines

Connecting Research Outputs

Community

Design Manual

DataCite Slide Deck

DataCite Brochure

DMP IDs

Introduction to Machine Actionable DMPs (maDMPs)

DataCite DMP IDs



DataCite Commons and PID Graph DataCite

Find Research with Data Cite Commons

DataCite Commons exposes:

the connections between DOIs in the form of citations, versions, and collections

the connections between content with DOIs, people (ORCID), research organizations (ROR), and funders (Crossref Funder ID) e.g. all works/funder or all works/organization

More information:

https://doi.org/10.5438/f4df-4817



Aggregation by Organization

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Publication Year

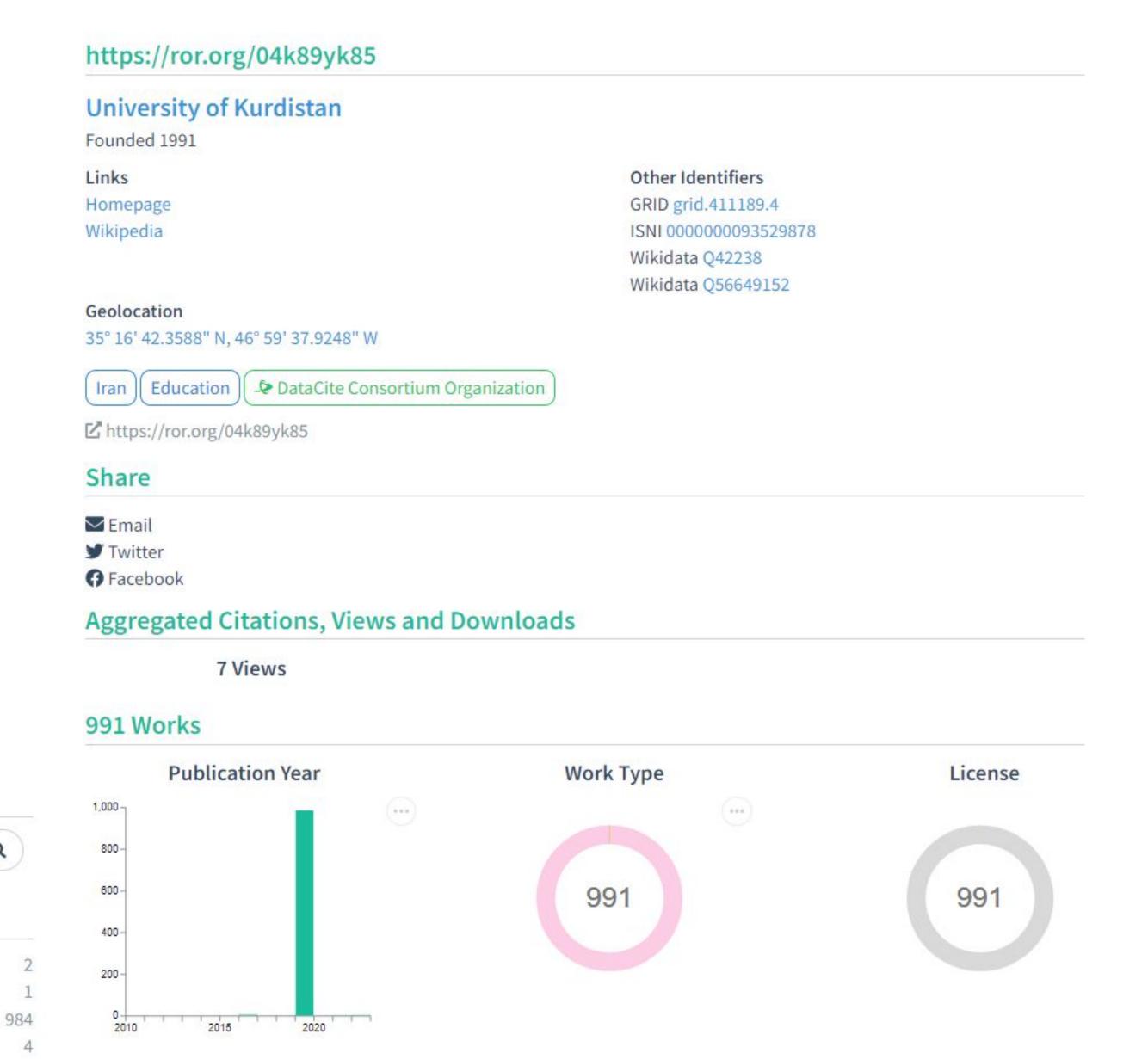
2022

2021

2019

2016

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GraphQLAPI



- ·Supports queries of the DataCite API using the GraphQL query language.
- •Contains: All Datacite DOIs ~9 million Crossref DOIs All ORCID iDs All ROR IDs All Crossref Funder IDs All re3data records
- •Support documentation (from DataCite): https://support.datacite.org/docs/datacite-graphql-api-guide
- •Video tutorial (from DataCite): https://www.youtube.com/watch?v=efvxGfU_oVM
- ·About GraphQL generally: https://graphql.org Jupyter notebooks demonstrating
- •FREYA use cases: https://github.com/datacite/notebooks
- •PID Forum for all PID Graph related questions: https://www.pidforum.org/c/pid-grap

DataCite Downstream Impact

Metadata is publicly available



Research Organization ————DataCite DOI Service

registers DOI with metadata

Metadata can be found/retrieved:

- 2.DataCite Commons
- ☐ 3.via our APIs
- 4..OAI-PMH

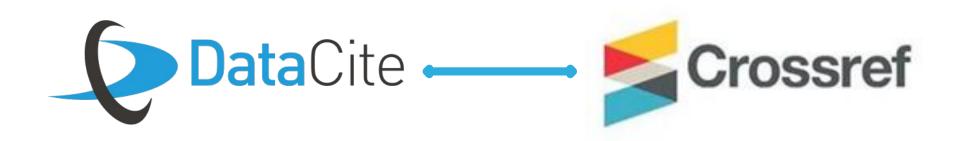
Metadata is harvested by:

- Mendeley Data
- ☐ Clarivate Web of Science
- Discovery Portals e.g. Base and OpenAIRE Explore
- Google Dataset Search
- Library Catalogs

DataCite Commons - Citations

License





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3,887 Citations 9,511 Views 2,383 Downloads

6,032 Works



https://doi.org/10.5061/dryad.2ht96

Data from: Relative accuracy of three common methods of parentage analysis in natural populations

Hugo B. Harrison, Pablo Saenz-Agudelo, Serge Planes, Michael L. Berumen & Geoffrey P. Jones Version 1 of Dataset published 2012 in DRYAD

Parentage studies and family reconstructions have become increasingly popular for investigating a range of evolutionary, ecological and behavioral processes in natural populations. However, a number of different assignment methods have emerged in common use, and the accuracy of each may differ in relation to the number of loci examined, allelic diversity, incomplete sampling of all candidate parents, and the presence of genotyping errors. Here we examine how these factors affect the accuracy of three popular parentage inference methods (COLONY, FaMoz and an exclusion-Bayes' theorem approach by Christie et al. (2010a)) to resolve true parent-offspring pairs using simulated data. Our findings demonstrate that accuracy increases with the number and diversity of loci. These were clearly the most important factors in obtaining accurate assignments explaining 75-90% of variance in overall accuracy across 60 simulated scenarios. Furthermore, the proportion of candidate parents sampled had a small but significant impact on the susceptibility of each method to either false positive or false negative assignments. Within the range of values simulated, COLONY outperformed FaMoz, which outperformed the exclusion-Bayes' theorem method. However, with 20 or more highly polymorphic loci, all methods could be applied with confidence. Our results show that for parentage inference in natural populations, careful consideration of the number and quality of markers will increase the accuracy of assignments and mitigate the effects of incomplete sampling of parental populations.

DOI registered October 24, 2012 via DataCite.



66 1 Citation ● 338 Views 🕹 141 Downloads

Dataset English

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Relative accuracy of three common methods of parentage analysis in natural populations

Hugo B. Harrison, Pablo Saenz-Agudelo, Serge Planes, Geoffrey P. Jones & Michael L. Berumen Journal Article published 2012 in Molecular Ecology

DOI registered via Crossref.

66 2 Citations

Journal Article

https://doi.org/10.1111/mec.12138



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