

Linking databases to code repositories with Throughput

March 3, 2020
WestGrid Virtual Presentation

Simon Goring

Kerstin Lehnert, Nick McKay, Steve Kuehn, Shane Loeffler, Anders Noren, Andrea Thomer, Socorro Dominguez



Introduction

Simon Goring

- Associate Scientist, Geography (UWisc)
- Adjunct Professor, Computer Science (UBC)

Motivated by a desire to help mobilize data to improve equity and reduce “time to science”.

Throughput (<https://throughputdb.com>)

A three year NSF Funded project through the EarthCube (<http://earthcube.org>) program to connect data resources in the geosciences and beyond.



Twitter: @sjGoring

Collaborators

Kerstin Lehnert - Columbia University EarthChem/SESAR/IGSNs

Nick McKay - Northern Arizona University (LinkedEarth)

Steve Kuehn - Concord University

Shane Loeffler - University of Minnesota (<https://flyovercountry.io/>)

Anders Noren - University of Minnesota (CSDCO, LacCore)

Andrea Thomer - University of Michigan School of Information

Socorro Dominguez - University of Wisconsin (UBC Data Science)

Interdisciplinary research is complicated.

Finding Appropriate Data -> Using Appropriate Methods -> Ensuring Reproducibility -> Obtaining Credit

Research is complicated.

Finding Appropriate Data -> Using Appropriate Methods -> Ensuring Reproducibility -> Obtaining Credit

Problems to Solve

1

Finding Appropriate Data

2

Using Appropriate Methods

3

Ensuring Reproducibility

4

Obtaining Credit

Problems to Solve

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Finding Appropriate Data

Knowledge of disciplinary archives (e.g., Neotoma, OpenContext).

2

Using Appropriate Methods

3

Ensuring Reproducibility

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Obtaining Credit

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Using Appropriate Methods

Time series analysis, data processing, R packages.

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Problems to Solve

1 Finding Appropriate Data

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Ensuring Reproducibility

Finding support, reproducing best practices, sustainability.

2 Using Appropriate Methods

Time series analysis, data processing, R packages.

Obtaining Credit

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Using Appropriate Methods

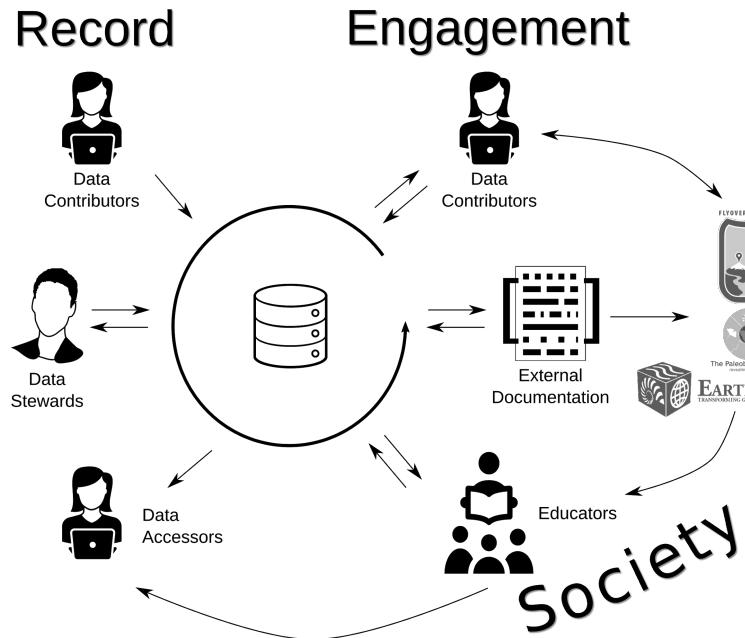
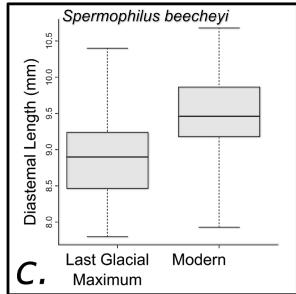
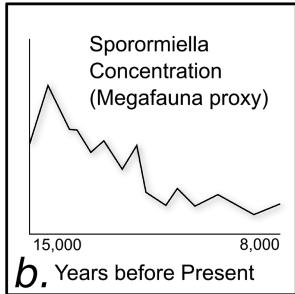
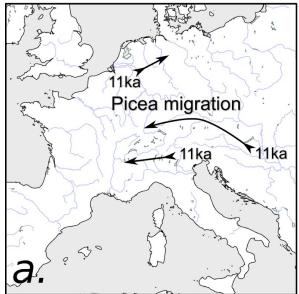
Time series analysis, data processing, R packages.

4

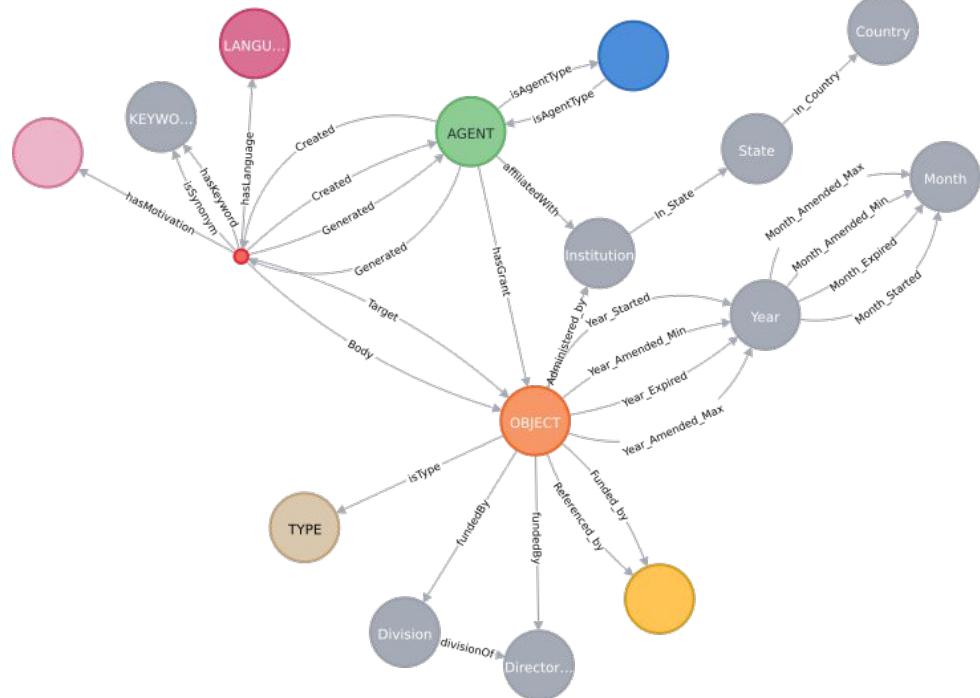
Obtaining Credit

Credit for methods and products beyond papers.

Mid-Scale Infrastructure



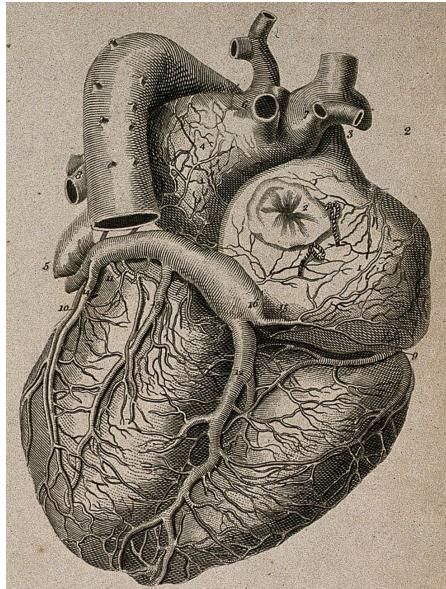
Solutions from a Graph



Throughput provides:

- A platform for data object annotation.
 - A tool to discover connections
 - A system for managing non-authoritative metadata
 - A tool to understand and track data and software citation

At its Heart, Annotations

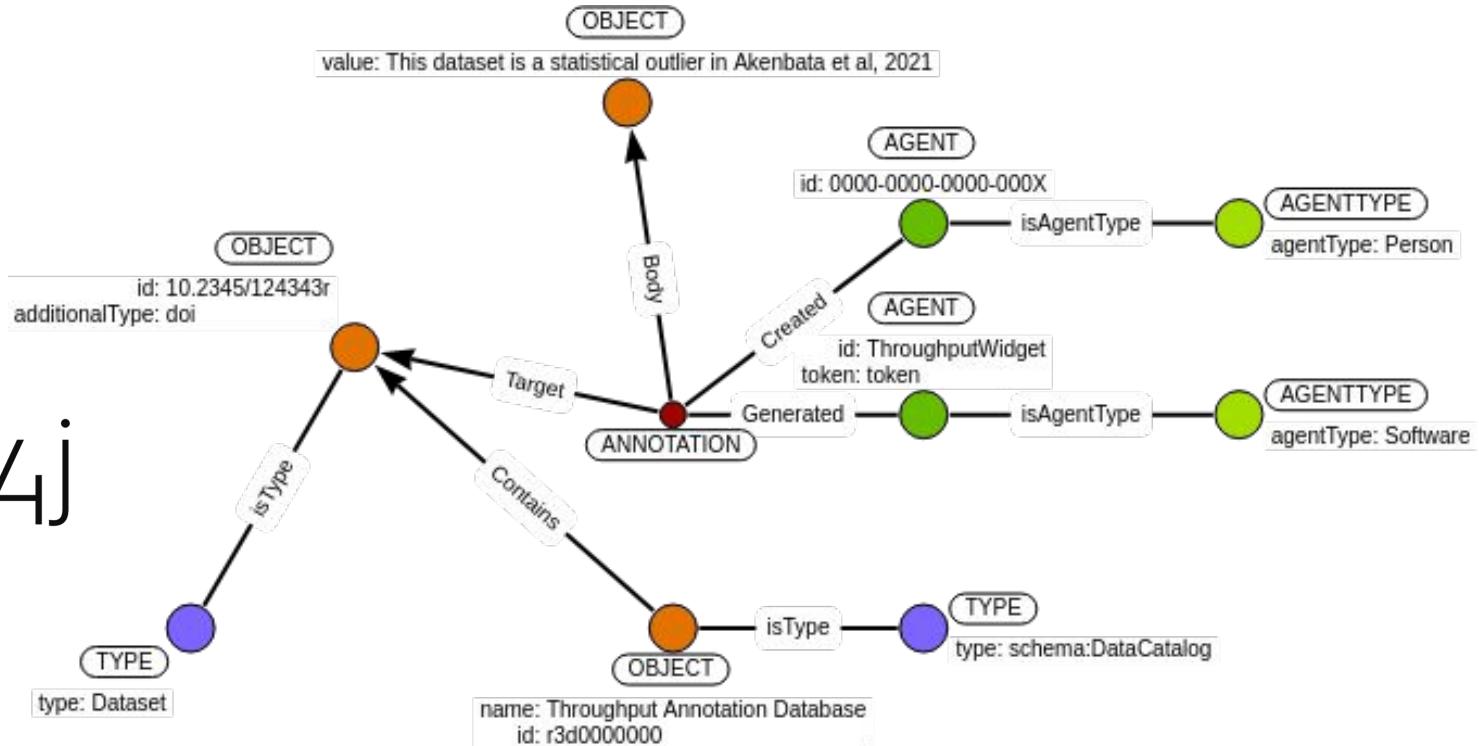


W3C Annotation model (<https://www.w3.org/TR/annotation-model>)

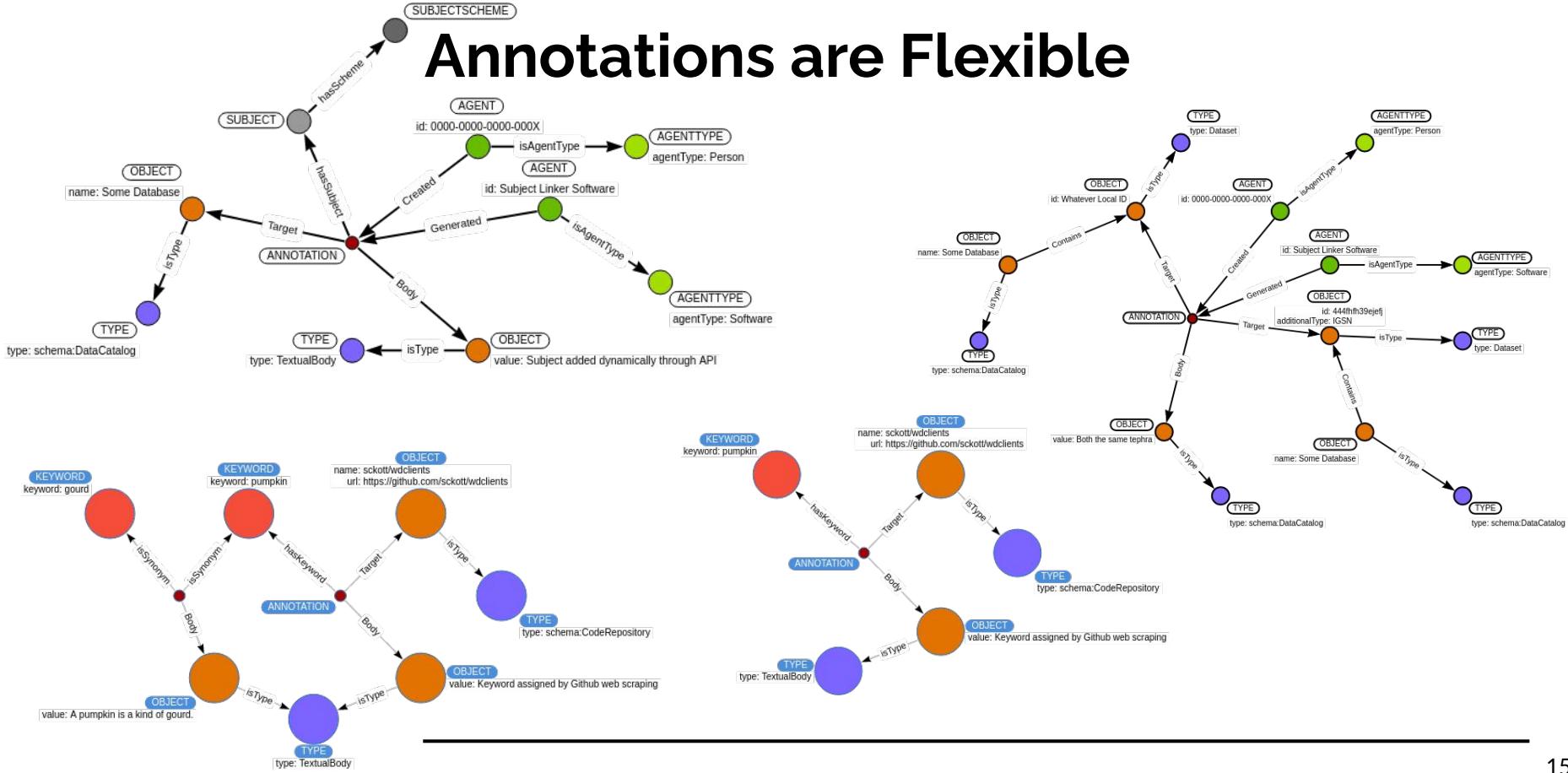
Extended using defined vocabularies (schema.org, DataCite, CrossRef, NSF)

- Annotations add information (“this dataset is cool!”)
- Annotations link records (“This GitHub repo is mentioned in this paper”)

At its Heart, Annotations

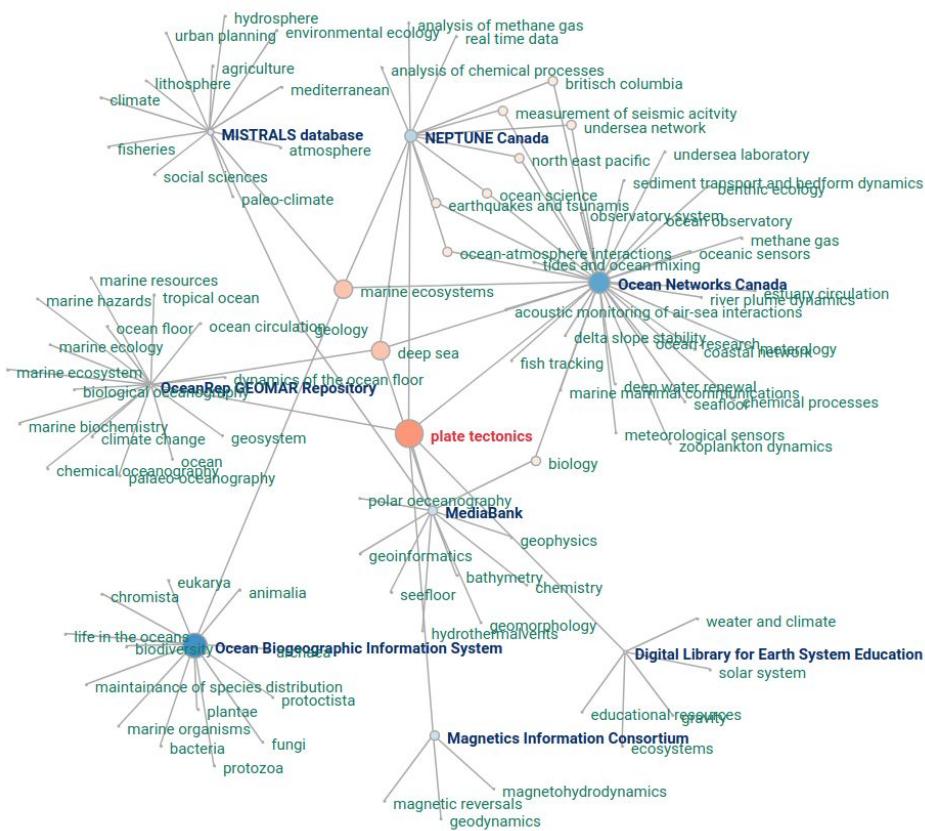


Annotations are Flexible



Linked Annotations Build Networks

- 300,000 Annotations
 - 43,000,000 object connections
 - 467,000 Scientific Grants
 - 160,000 Research Articles
 - 2,500 Databases
 - 74,000 Code Repositories



Networks for Knowledge

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Knowledge of disciplinary archives (e.g., Neotoma, OpenContext).

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Time series analysis, data processing, R packages.

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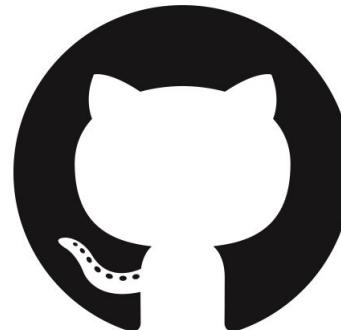
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Time series analysis, data processing, R packages.



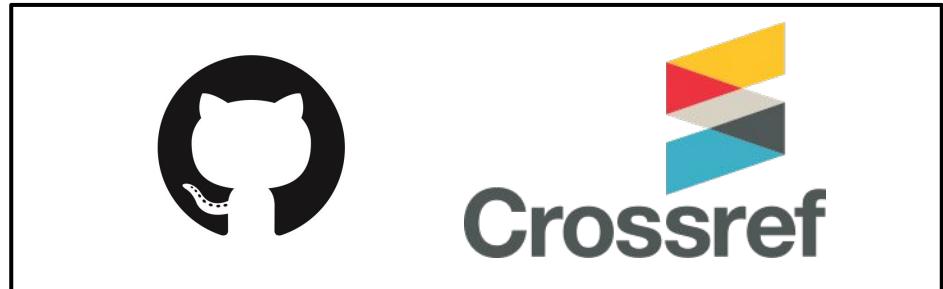
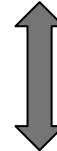
Networks for Knowledge

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Discovering Links

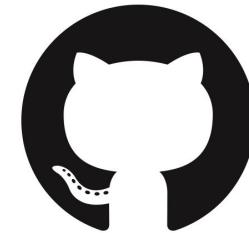
AdamWilsonLabEDU/geo503-2018-finalproject-nathandubinin
final_project.R

```
 9 library(grid)
10 library(neotoma)
11 library(analogue)
12
13 # Info from Neotoma Explorer (Data originally from European Pollen Database)
...
31 # Getting data using Neotoma's API for
32 # Lake Sambösön
33 sam <- get_site(sitename = 'Lake Sambösön')
34 sam_pollen<-get_dataset(sam)
```

R Showing the top six matches Last indexed on 14 Dec 2018



Neotoma
R package



xDeepDive (geoDeepDive)

13,415,017 documents



47,417 added this month

14,140 added this week

2,539 added in the last 24 hours

<https://geodeepdive.org/>



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xDeepDive

Search & Tagging with:

- Full Text
 - Regular Expression
 - NLP Models



 Crossref



Shanan E. Peters Dean L. Morgridge Professor

Sedimentary Geology and Paleobiology,
GeoInformatics



EARTH CUBE

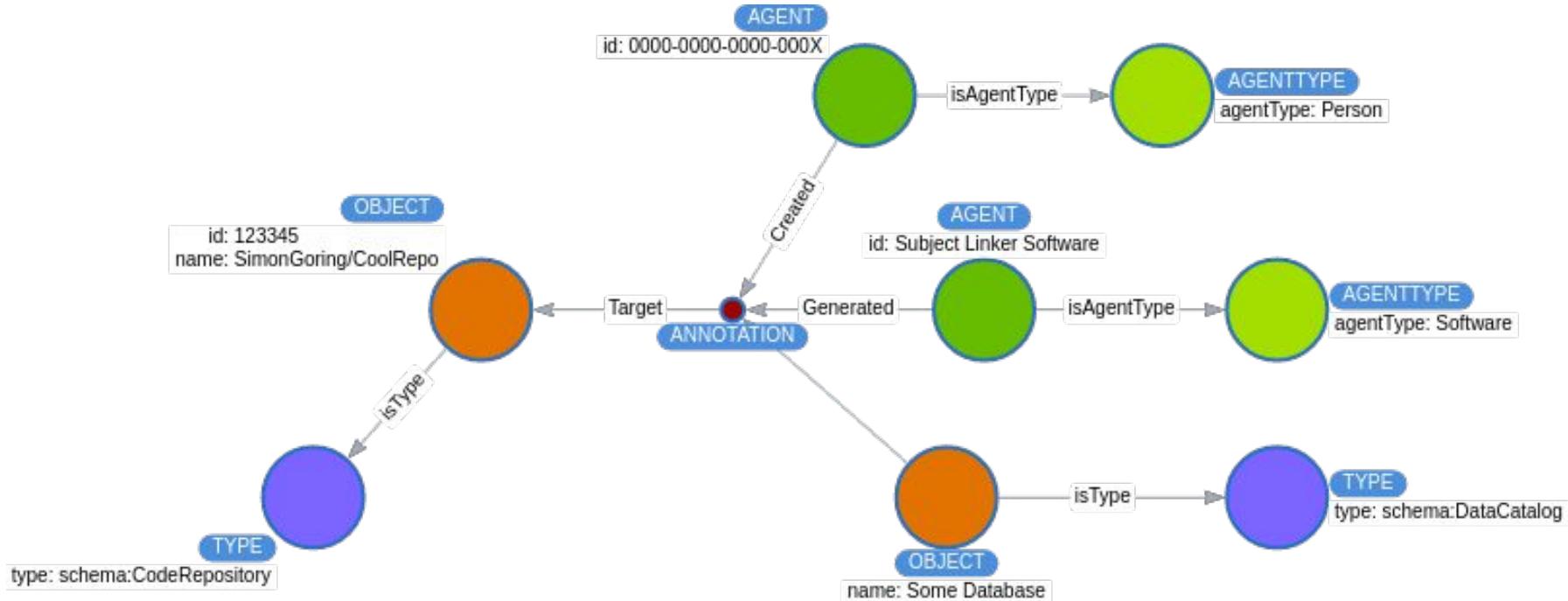
TRANSFORMING GEOSCIENCES RESEARCH



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DB to GitHub Links



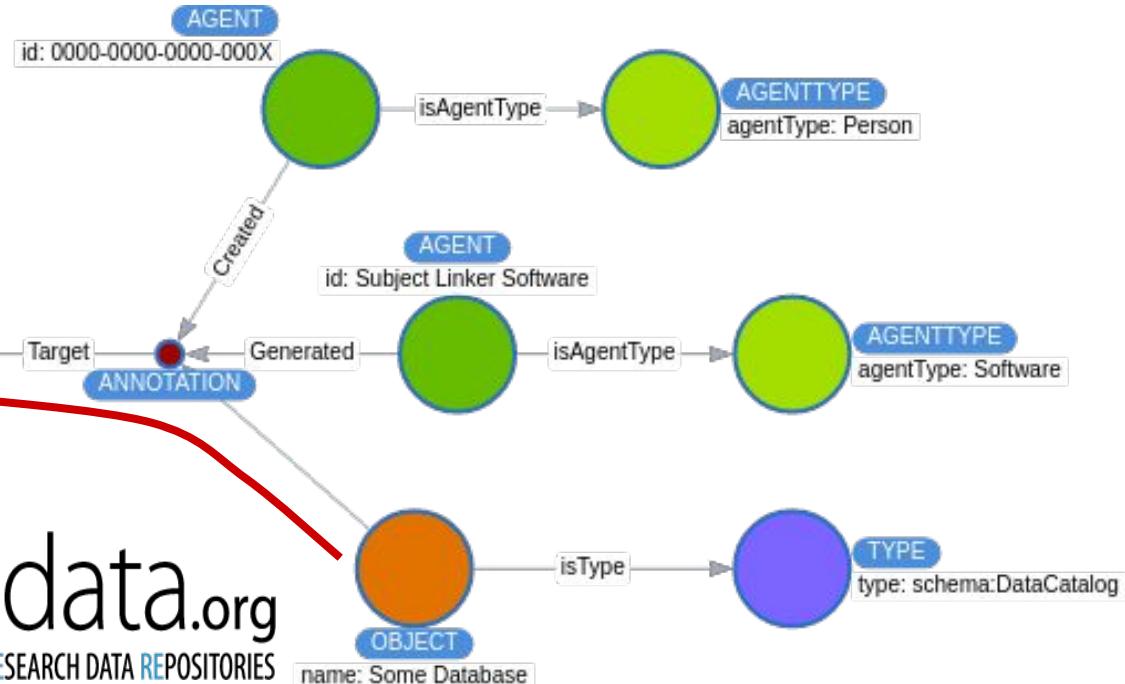
DB to GitHub Links

Use of External APIs and
PIPs as a core element of
the Throughput Database



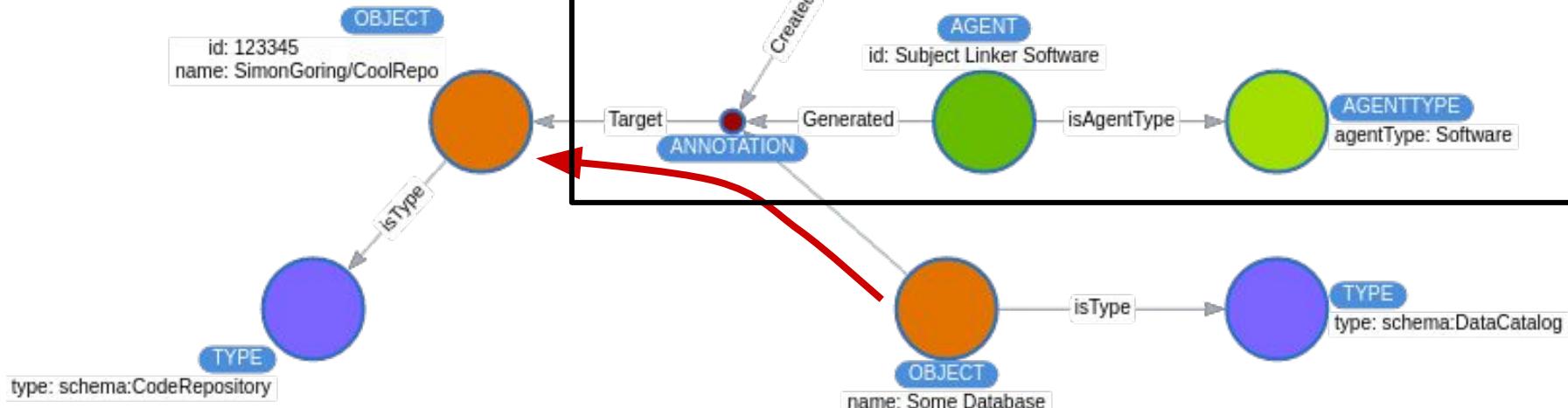
OBJECT
id: 123345
name: SimonGoring/CoolRepo

TYPE
type: schema:CodeRepository



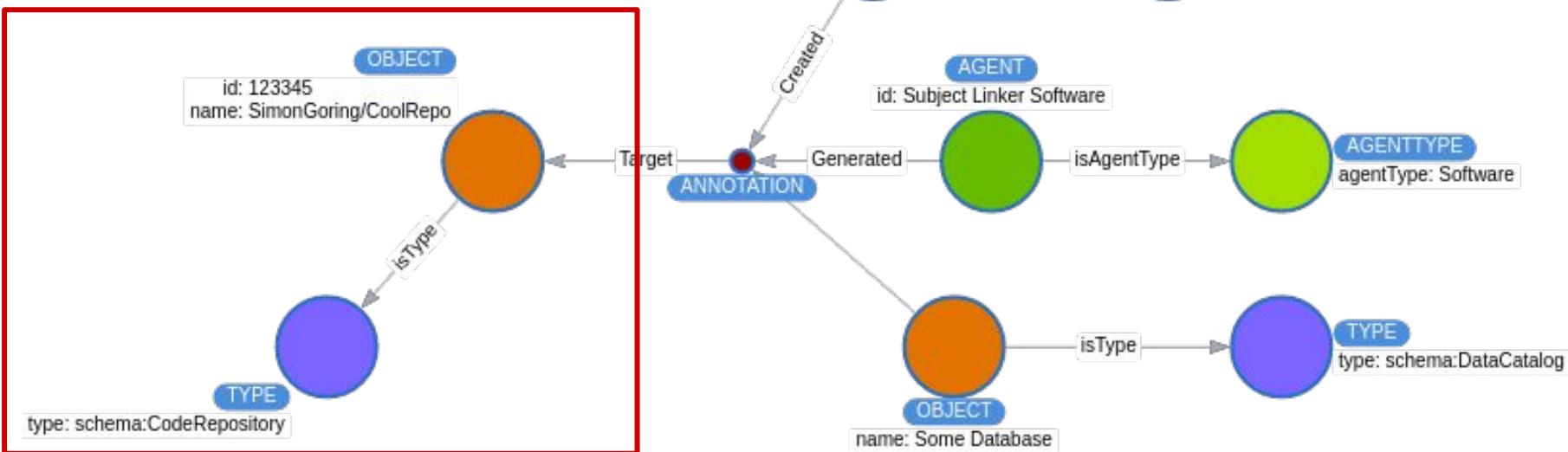
DB to GitHub Links

Content to manage
Trust and Provenance

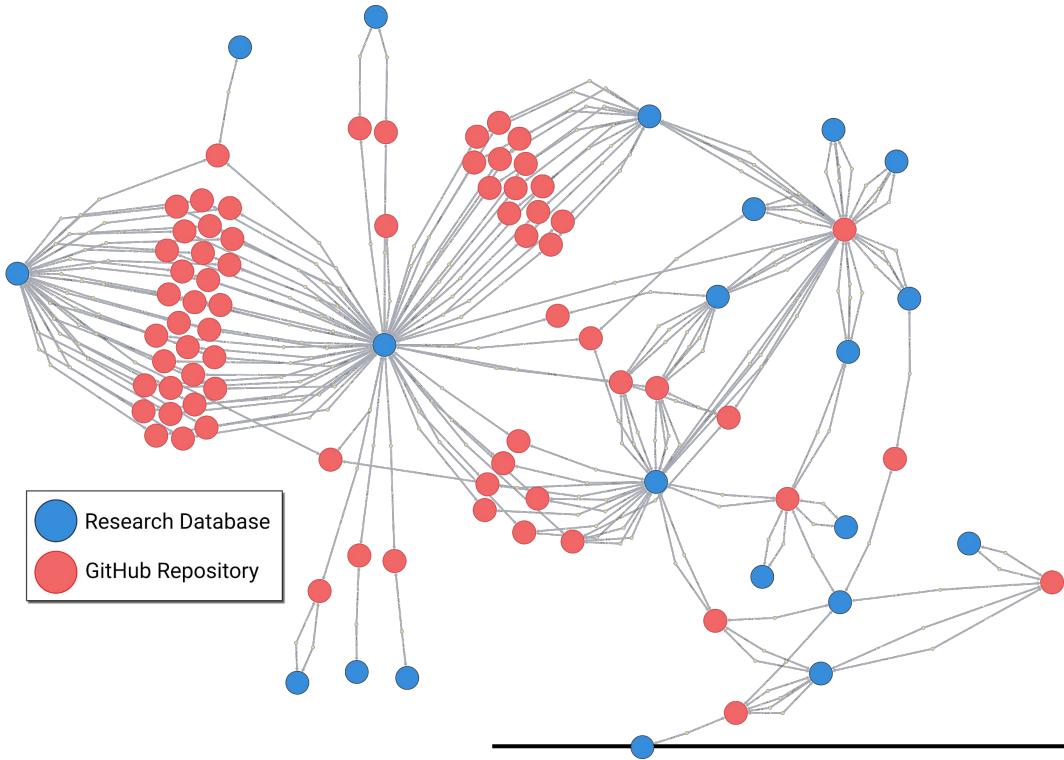


Model Is Extensible

TYPEs include Journal Articles,
Datasets, Grants, Text
Annotations and others.

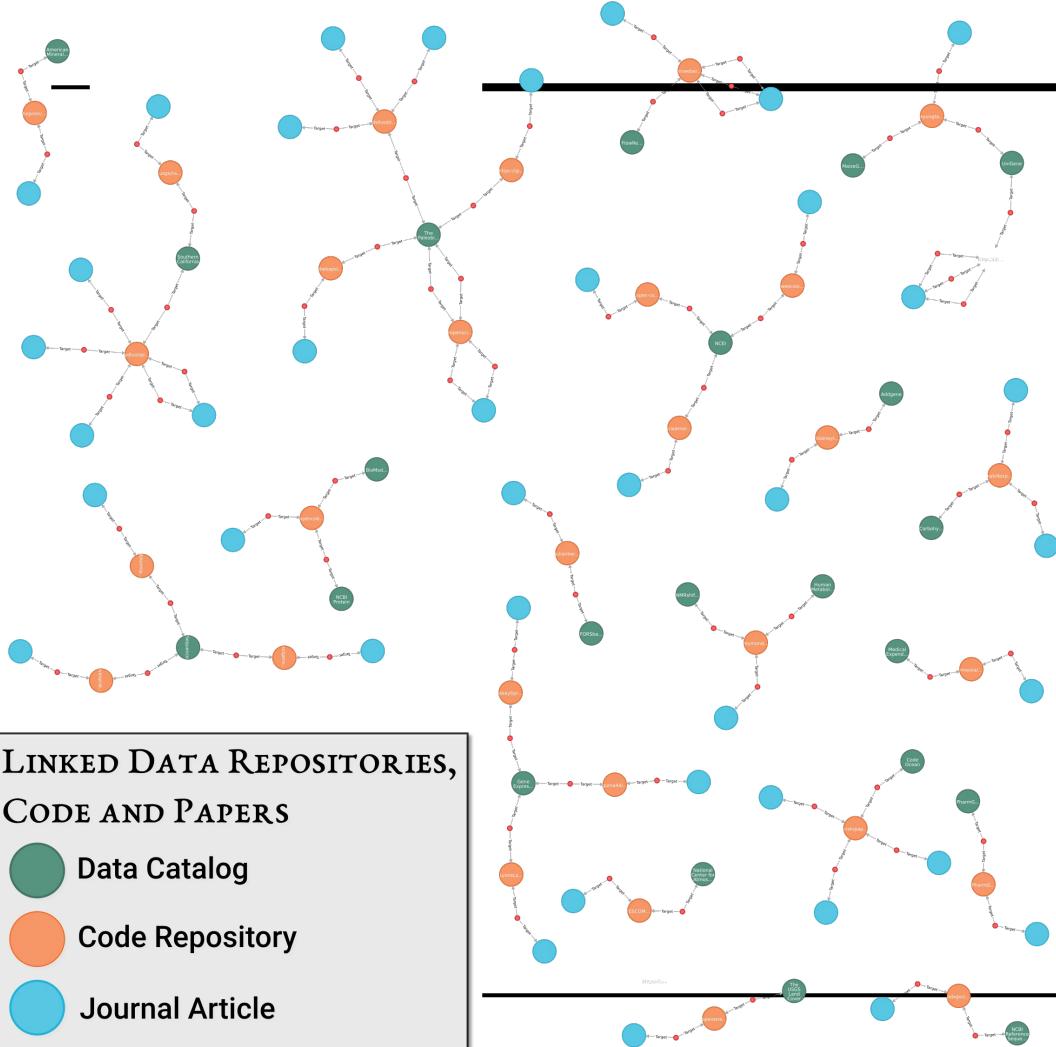


Links Across Data Resources



Patterns of data alignment

Variety often poses one of the greatest challenges in Big Data analysis. Are these repositories sources of information on how best to undertake data alignment?



Data Discovery

Tracking Data Use

Understanding Best Practices

Retrospective Credit & Linking

LINKED DATA REPOSITORIES,
CODE AND PAPERS

Data Catalog

Code Repository

Journal Article

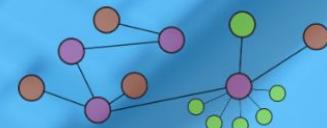
Data Discovery & Use

[Home](#)

[About Throughput](#)

[GitHub Code](#)

Throughput Data and Software
Annotation



Annotation Editor

Challenges In Metadata Recovery

README Files (<https://github.com/throughput-ec/Template>)

Proper citation & identification of source data

Optical Character Recognition issues in legacy publications

Lack of standards for Jupyter/RMarkdown documents

(<https://github.com/earthcube/NotebookTemplates>)

Challenges in Code Discovery

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Repository Variety

GitHub Repository Typology

Throughput

Educational: Elements of this repository serve to house educational and instructional resources. This may include: Books, assignments, class notes, and tutorials. This also includes manuals and training materials for the use of software. This does NOT include the results of class projects.

Analysis: This repository contains the data and code used as a primary analysis for some sort of research project. These are custom data analysis pipelines, not meant to be generally reusable. This category includes use and reuse of data. This includes: comparative reuses, original research studies, meta-analyses, statistical method development, and reproducibility tests.

Software development: Elements of this repository serve to build freestanding tools of any sort including libraries, plugins, frameworks, etc. This includes use of data to pilot tools. Includes calls to database API or library or other computational interface. Includes calls to R libraries

Storage: Repository stores copies of data from research databases.

Miscellaneous: This category is for repositories that are out of scope of this typology. This includes:

Scraping database registries: Repository contains lists of research databases and associated metadata.

Articles referencing database: Repository contains articles that link to research databases.

Informational link to database: Links to research database's homepage or another informational page. Not to a dataset.

Can't categorize/not enough information

Github repository no longer accessible: Github repository gives a 404 error.

Research database no longer accessible: Research database linked from GitHub repository gives a 404 error.

ML Classification and Tagging

The screenshot shows a user interface for managing biological resources. At the top, there are two tabs: "Data Catalogs" (selected) and "Code Repositories". Below the tabs are two buttons: "Get Citations" and "Show Unselected Resources".

bio-tools/content

Drop

Pathway Commons, Complete Genomics, EMBL-EBI, ArrayExpress, MaizeGDB, DrugBank, OMICtools, NITRC, MEROPS, Gramene, PLEXdb, Biosamples, Golm Metabolome Database, REBASE, GenomeRNAi, AmoebaDB, FlyBase, NCBI Epigenomics, Rat Genome Database

Experimental repo of bio.tools content for augmentation with other sources of tool information

BioContainers/tools-metadata

Drop

Pathway Commons, Kyoto Encyclopedia of Genes and Genomes, Complete Genomics, EMBL-EBI, ArrayExpress, MaizeGDB, NITRC, MEROPS, Gramene, PLEXdb, Biosamples, Golm Metabolome Database, REBASE, GenomeRNAi, AmoebaDB, FlyBase, NCBI Epigenomics, Rat Genome Database

Repository to storage the tools metadata.

ML Classification and Tagging

The screenshot shows a user interface for managing biological resources, likely a GitHub repository or a similar platform. At the top, there are tabs for "Data Catalogs" and "Code Repositories". Below these are buttons for "Get Citations" and "Show Unselected Resources". On the left, there is a GitHub logo icon.

bio-tools/content

This section lists various databases and tools: Pathway Commons, Complete Genomics, EMBL-EBI, ArrayExpress, MaizeGDB, DrugBank, OMICtools, NITRC, MEROPS, Gramene, PLEXdb, Biosamples, Golm Metabolome Database, REBASE, GenomeRNAi, AmoebaDB, FlyBase, NCBI Epigenomics, Rat Genome Database. It is described as an "Experimental repo of bio.tools content for augmentation with other sources of tool information".

BioContainers/tools-metadata

This section lists the same set of databases and tools as the first section. It is described as a "Repository to storage the tools metadata".

Each resource entry has three status indicators: a green square with a question mark, a red square with a drop icon, and a blue square with an exclamation mark.

Active Development & Research

Implementing end-user annotation widget for data resources

Characterizing patterns of data use and reuse in the
geosciences

Feature extraction & metadata generation from publication

Data repository recommendation system

Continued integration of legacy & informal knowledge into
the research data graph

Active Development & Research

Annotations

5

Implementing end-user annotation widget for data resources



Throughput API

Characterizing patterns of data use and reuse in the geosciences

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Active Development & Research

Implementing end-user annotation widget for data resources

**Characterizing patterns of data use and reuse in the
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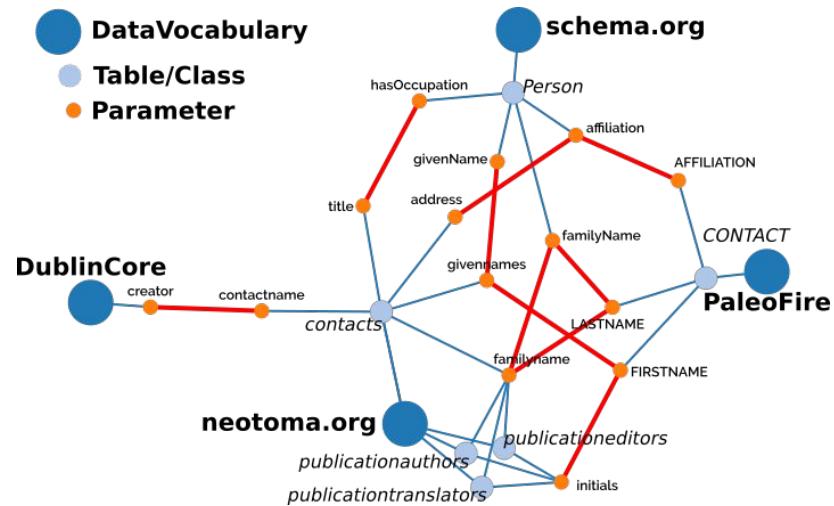
Data repository recommendation system

Continued integration of legacy & informal knowledge into
the research data graph

Characterizing patterns of data use

Volcanic Eruptions and Their Repose, Unrest, Precursors, and Timing

National Academies of Sciences, Engineering, and Medicine 2017. Washington, DC: The National Academies Press. <https://doi.org/10.17226/24650>



Active Development & Research

Implementing end-user annotation widget for data resources

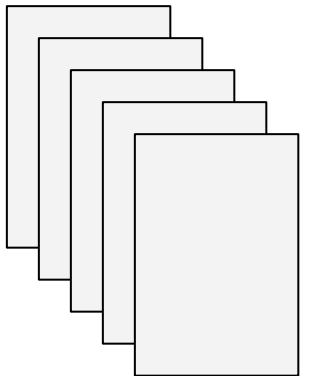
Characterizing patterns of data use and reuse in the
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Data Pipeline



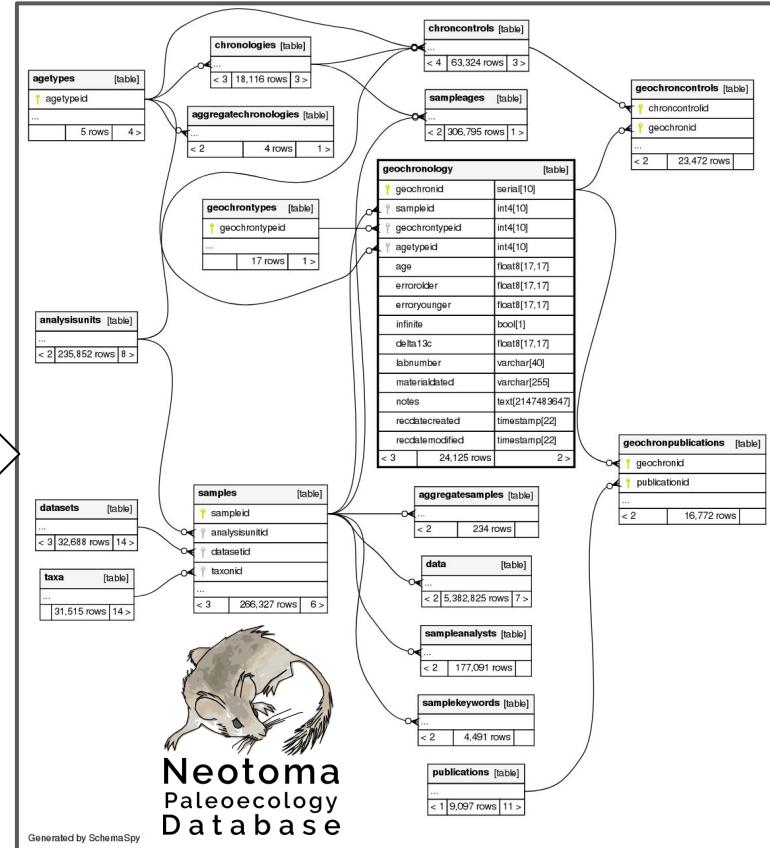
regex
tagging

ML
Prediction



Socorro Dominguez

bit.ly/emptyspacegdd



Active Development & Research

Implementing end-user annotation widget for data resources

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geosciences

Feature extraction & metadata generation from publication

Data repository recommendation system

**Continued integration of legacy & informal knowledge into
the research data graph**

Thanks!

Technical Notes

1. License - MIT (<http://github.com/throughput-ec/throughputdb>)
2. Data Standards Implemented
 - a. W3C Annotation (https://github.com/throughput-ec/throughputdb/tree/master/cypher_anno_examples)
 - b. Schema.org
 - c. existing (parent) metadata formats
3. Notebooks
 - a. DeepDive work using Jupyter notebooks & Dockerization (<https://github.com/throughput-ec/UnacquiredSites>)
 - b. DB/API/App all reproducible (<http://github.com/throughput-ec>), Dockerization to follow.
4. Fully version controlled (all work at <http://github.com/throughput-ec>)
5. Services
 - a. REST API using JSON (JSON-LD to follow), Documented using OpenAPI v3.0 (developed but buggy: <https://throughputdb.com/api-docs>)
6. Data
 - a. Snapshots deposited in Figshare (<http://doi.org/10.6084/m9.figshare.12731138>)