



The Genome Citation Service:

Capturing JGI data citations for
comprehensive impact assessment

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JGI: A DOE SC User Facility

- Part of the **Lawrence Berkeley National Laboratory**
- Located in **Berkeley, California**
- **~400 Staff**, including graduate students and postdocs
- **2,000+** Active PIs/collaborators per year
- **10,000+** Online users per year



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JGI User Programs

SCIENTIFIC PEER REVIEW

40%
Community
Science Program
(CSP)

10%
Facilities
Integrating
Collaborations for
User Science
(FICUS)



30%
Bioenergy
Research Centers
(BRC)

10%
Director's Science

10%
Biological and
Environmental Research
Support Science (BERSS)

Plant



Fungal, Algal



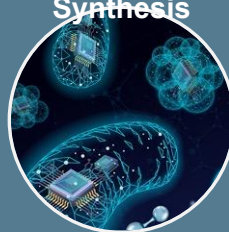
Metagenome



Microbial



DNA
Synthesis



Metabolomics



2° Metabolites



JGI Public Resources



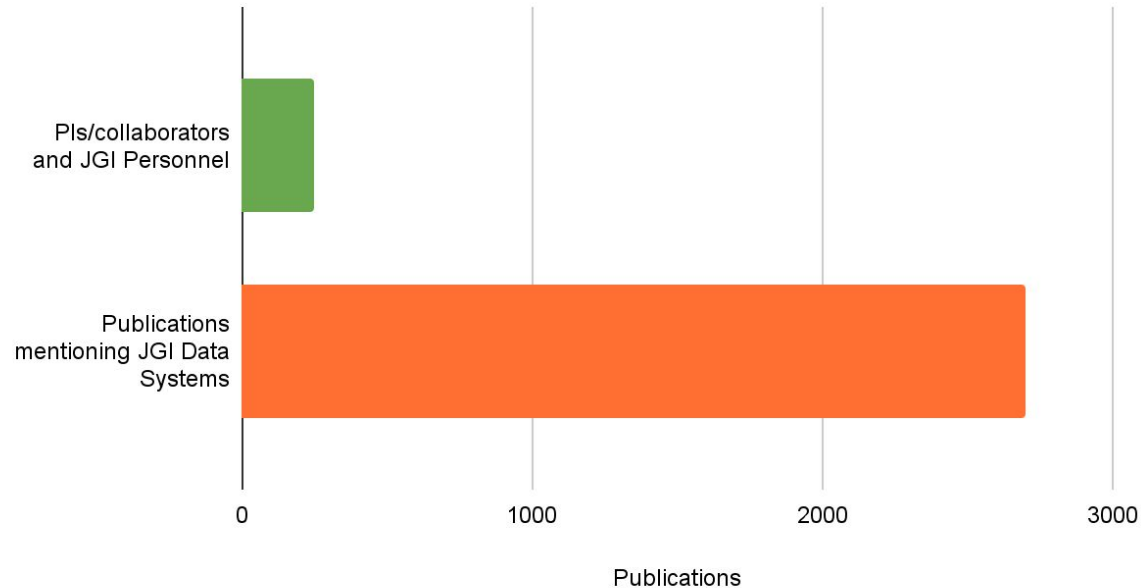
JGI Systems



External Systems

How do JGI user groups compare?

FY21 Publications



Raw data source:
 Dimensions

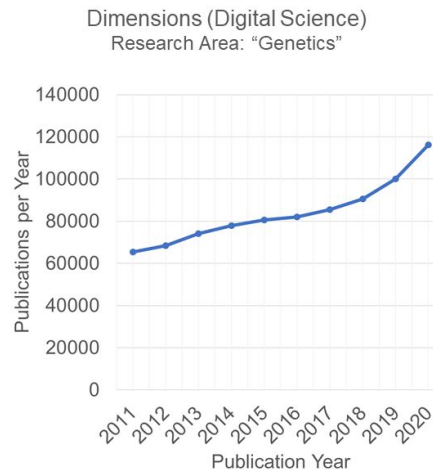
Scalability: Publications, Data, & Metadata

- Large amounts of data
- Rich and diverse metadata
- Growing body of literature

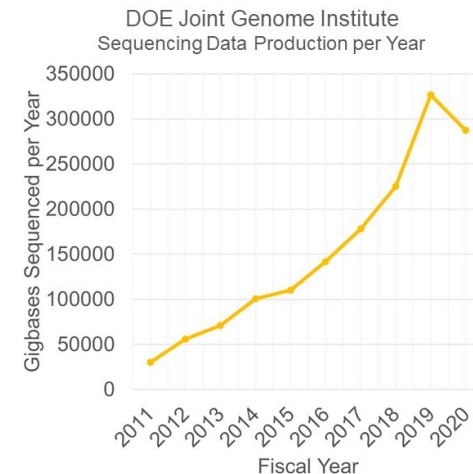


- *Too many citations* –
- *Using too many means* –
- *Of too many products* –

To identify manually



Net increase in yearly output: 2011-2020: 77.53%
Publication total 2011-2020: 840,780

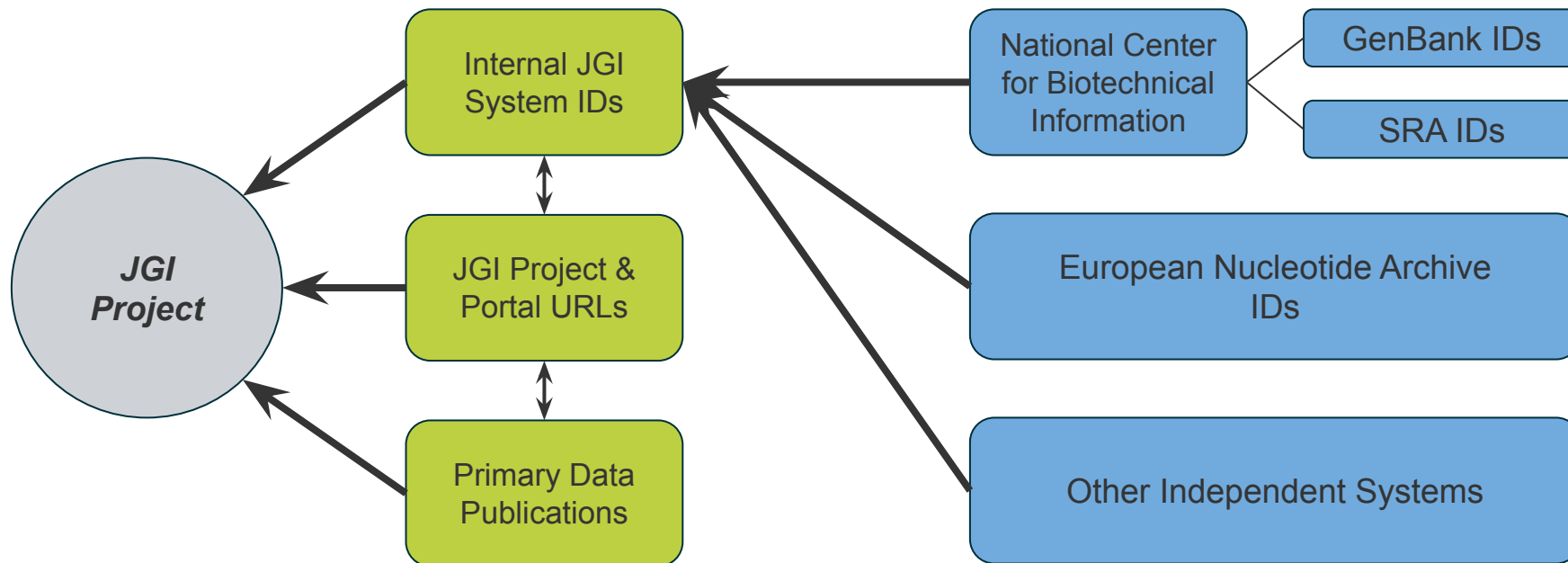


*FY2020 sequencing output affected by the COVID-19 pandemic.

Raw data source:

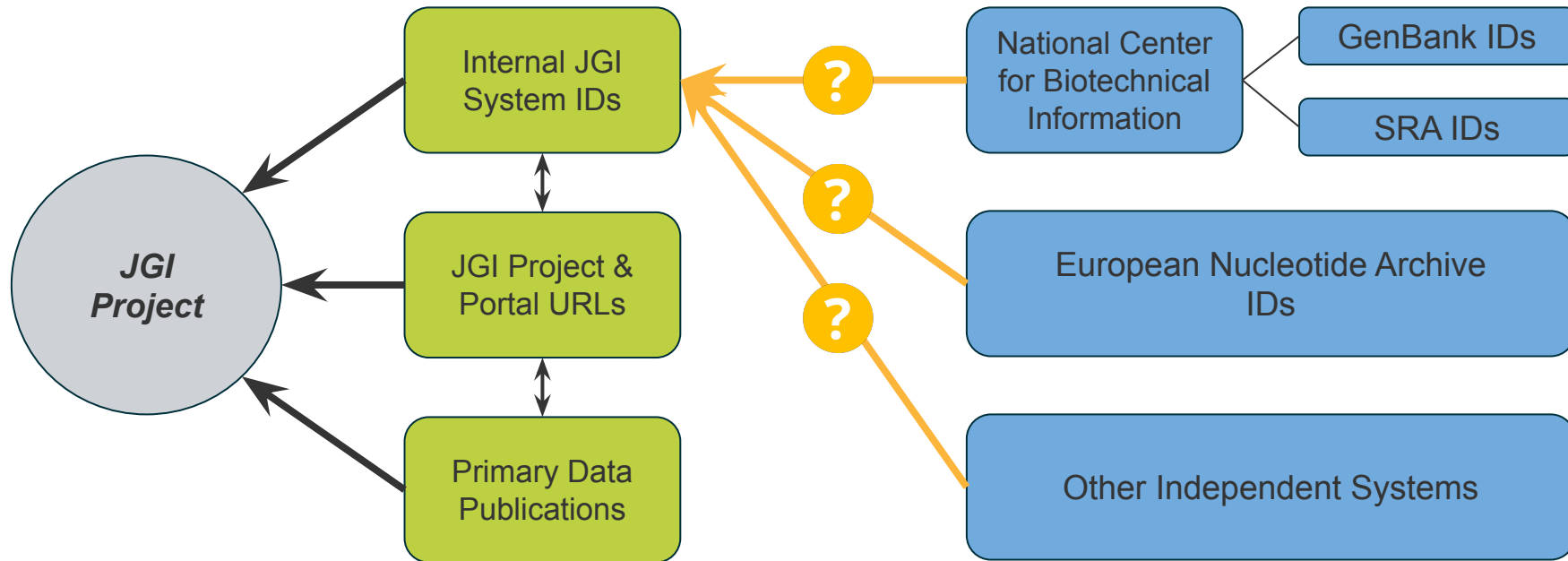


Incomplete Metadata



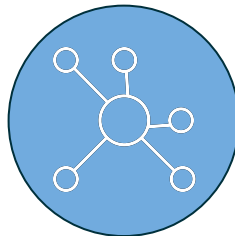
The Problem

Is a cited identifier linked to JGI? What project(s) is it linked to?



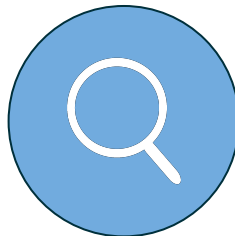
Automation: The Genome Citation Service (GCS)

Problem:
Incomplete Metadata



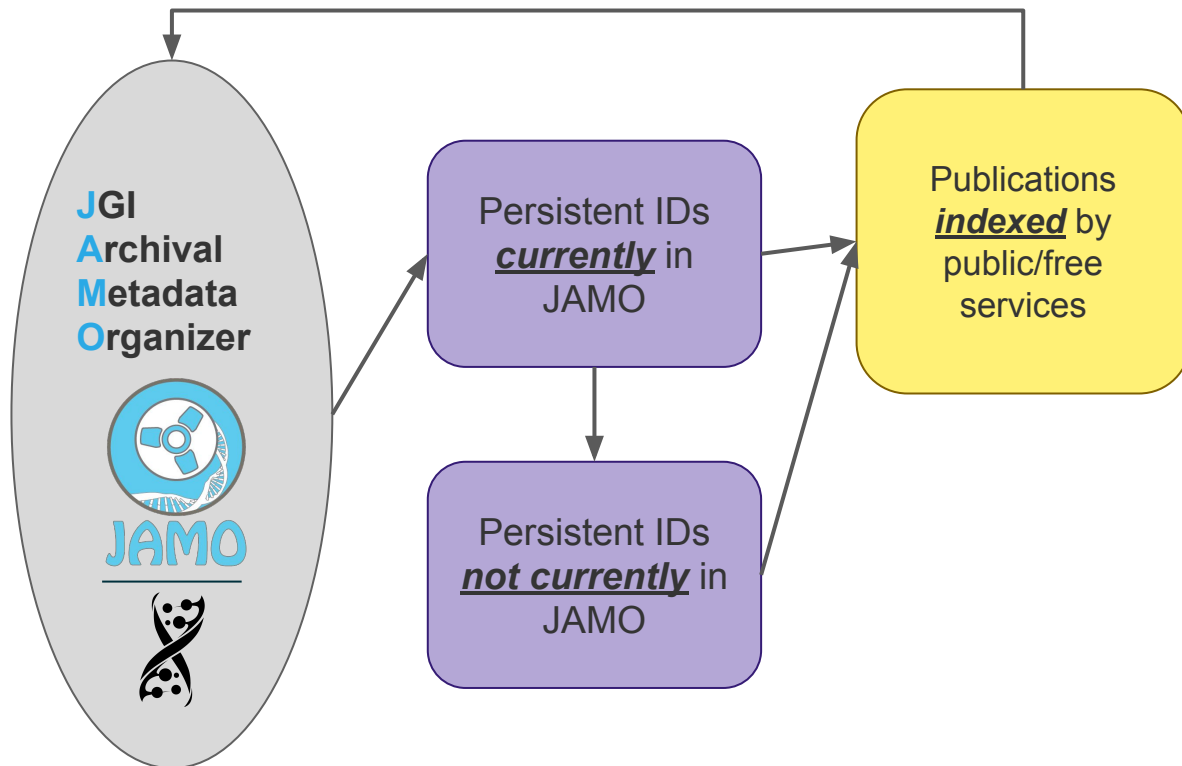
Solution:
Automatic, incremental discovery and traversal of additional linked data resources

Problem:
Scalability



Solution:
Automatic queries of public literature and cataloging of pathways back to JGI data resources

The Genome Citation Service

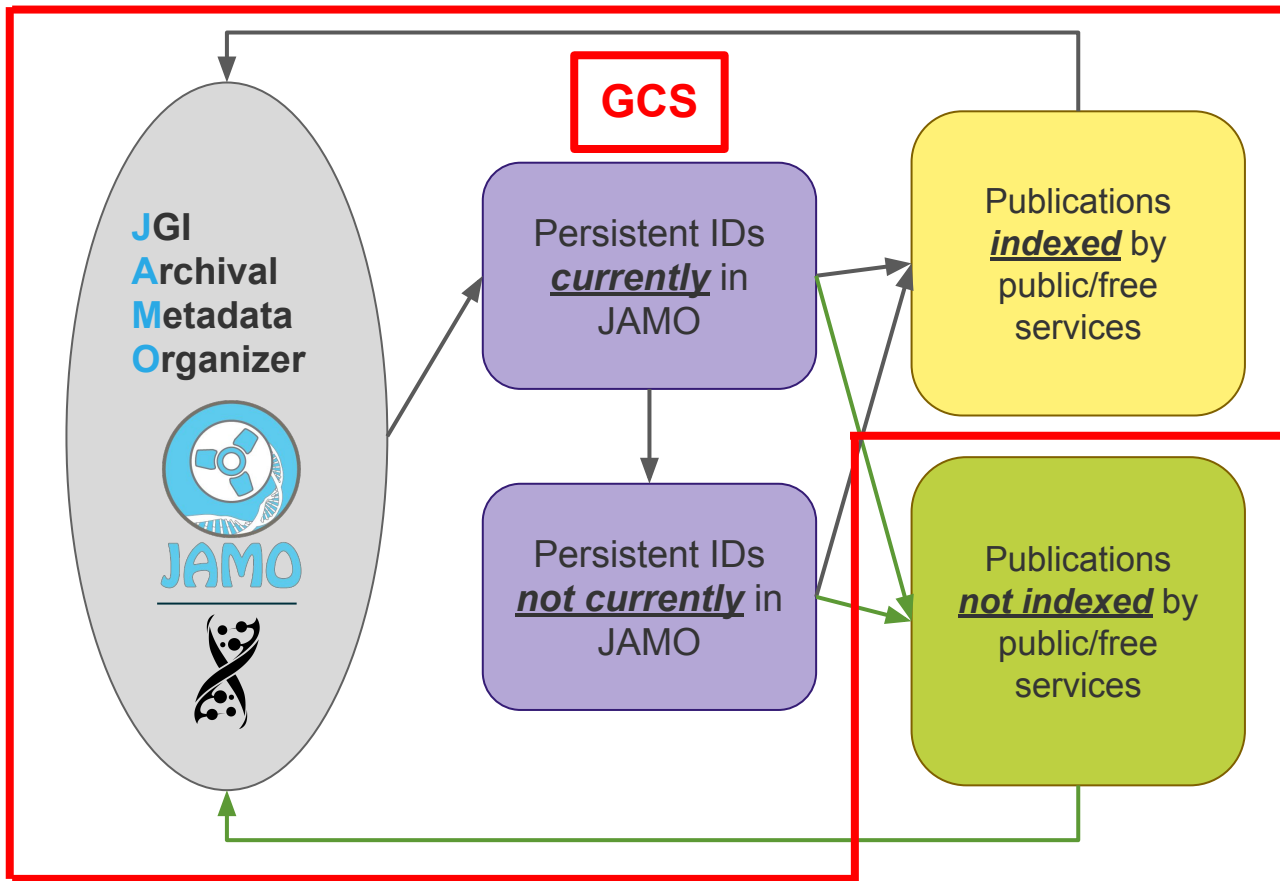


PubMed



PubMed
Central

The Genome Citation Service



PubMed

PubMed Central

Dimensions

Evaluating the GCS

How well did the GCS perform?

- **Methodology**

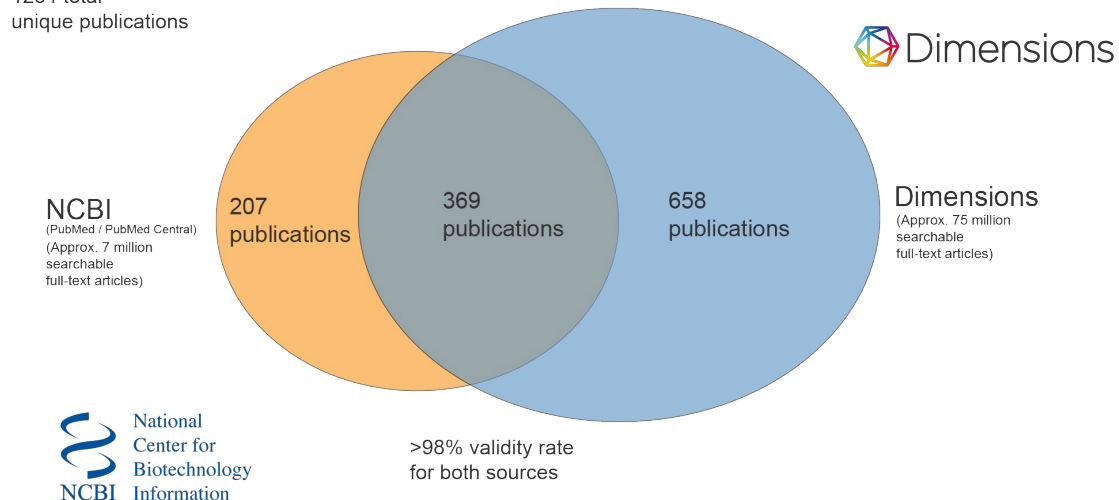
- 300 JAMO records (stratified, random)
- Search public and subscription sources for linked publications
- Manually evaluate hits

- **Results**

- >98% Precision
- 1234 total publications
- 90% previously unidentified

Publications identified through the NamesforLife / JGI "Genome Citation Service" using two distinct publication databases and a stratified random sample of 300 JAMO records

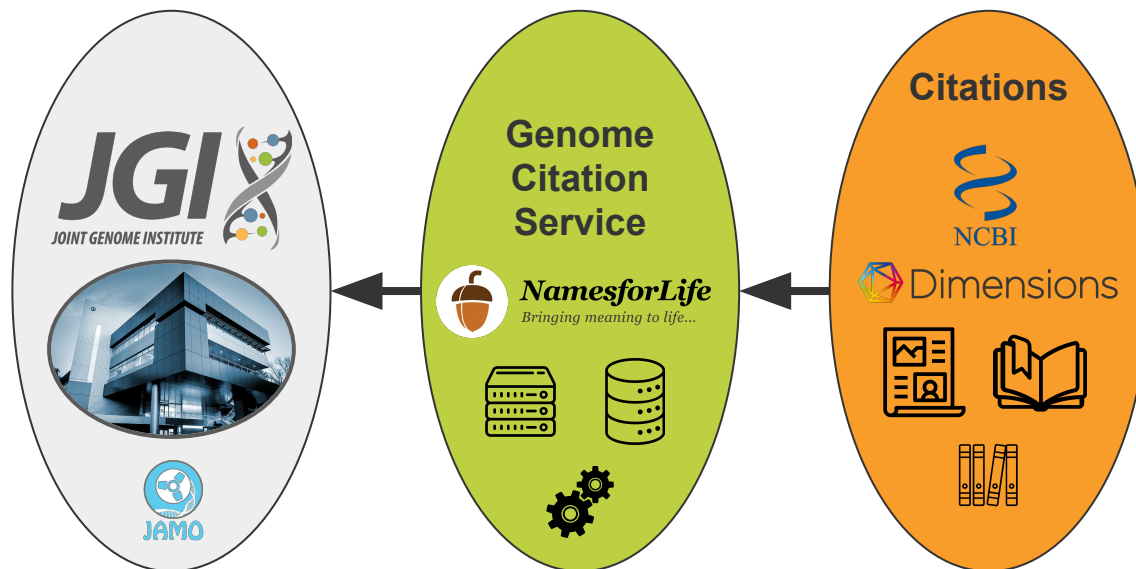
1234 total
unique publications



GCS Results: Implications

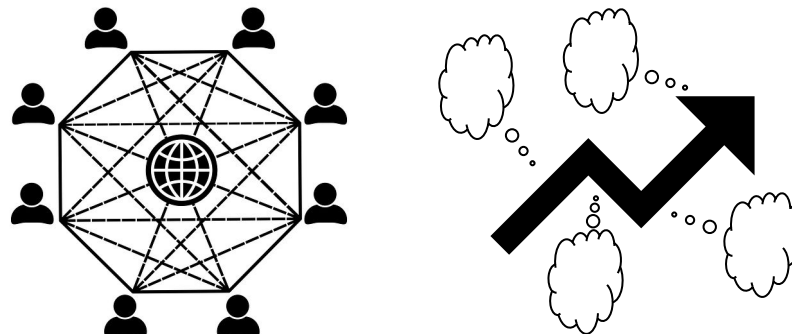
How does JGI benefit from the Genome Citation Service?

1. Thousands of JGI data citations now potentially identifiable
2. Little or no manual effort required

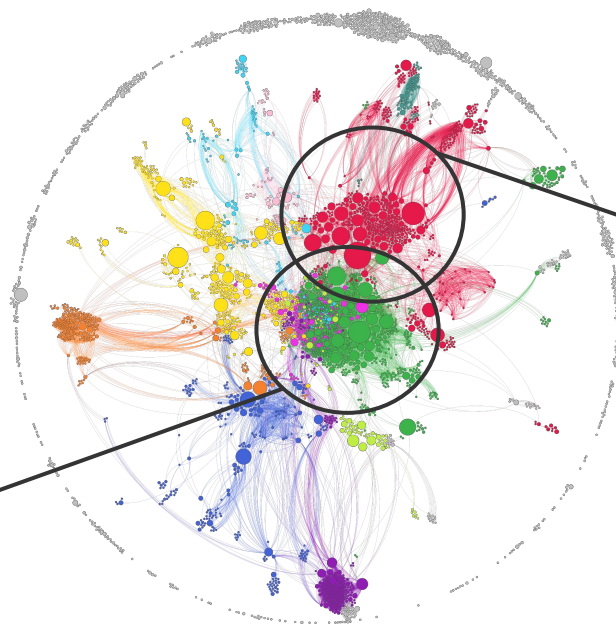


How can we use this information for impact assessment?

- **Community Analysis**
 - Coauthor and JGI user networks
- **Topic Analysis**
 - Data use trends
 - DOE goal alignment
- **New researcher metrics**
 - Equitable credit attribution



Which PI/Collaborator groups are producing data of scientific interest?



Group B

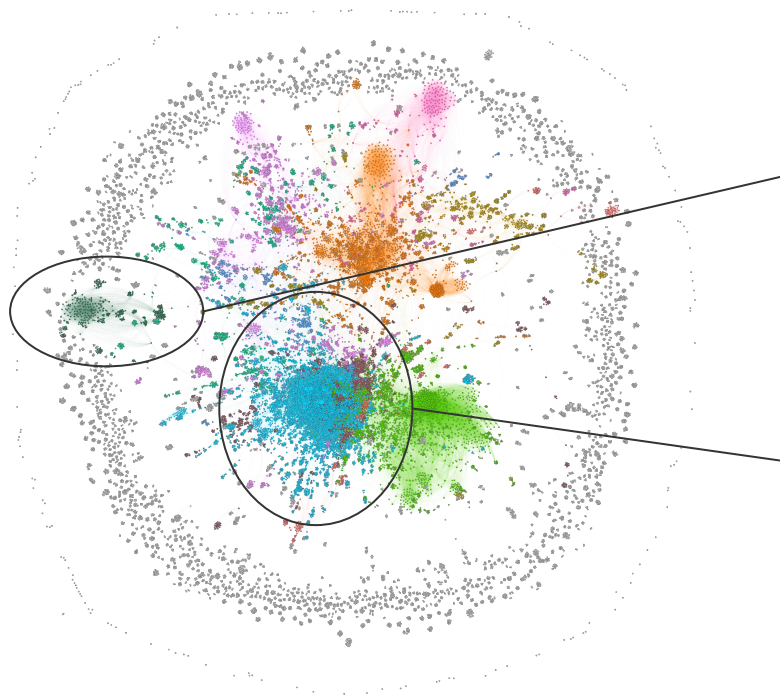
- X proposals with
- Y datasets receiving
- Z total data citations
- Contains many contributors to JGI's 1000 Fungal Genomes Project

Group A

- X proposals with
- Y datasets receiving
- Z total data citations
- Consists largely of DOE Bioenergy Research Center (BRC) personnel

JGI Proposal Contributors

Can we identify distinct communities among users of public JGI resources?



Blue Group* vs. Teal Group

- Total Authors:
 - 5015
 - 378
- Top Institutions:
 - DOE Labs, American & European Universities
 - European universities
- Top concepts:
 - Evolution, Pathogens, Biomass degradation
 - Bioinformatics, Ontology
- Publications
 - 1774 publications (~18 average citations)
 - 123 publications (~30 average citations)

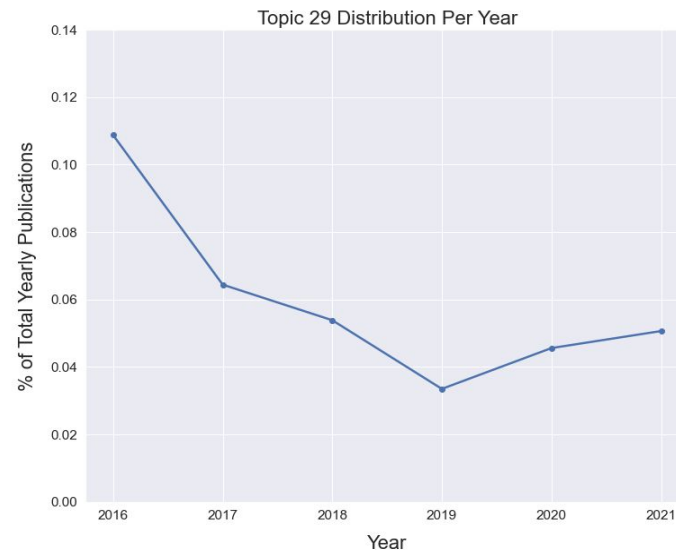
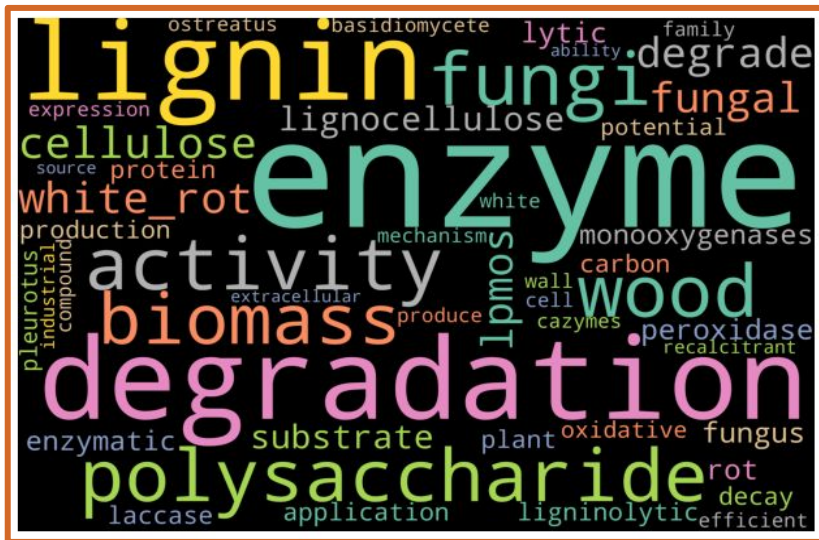
JGI Users
(mock subset)

Raw data source:
 Dimensions

*Top authors in Blue Group are JGI PIs & Collaborators

Topic & Concept Analysis

What 'topics' are trending amongst our user base?



This 'topic' seems to be losing prominence among citing publications. It consists of 341 publications from our test set, and is decreasing in total publication share over time.

Topic & Concept Analysis

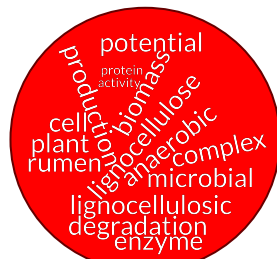
Are there counter-intuitive use cases for JGI data?

JGI Proposal: "Transcriptomic Characterization of Anaerobic Gut Fungi", 2014 549 Downstream Citations

Proposal Text:

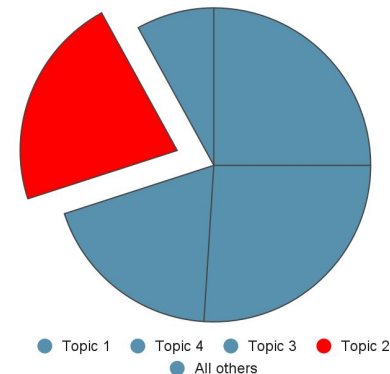
"The overall goal of our efforts is to develop the tools to engineer anaerobic gut fungi as novel platform organisms for **biofuel** production from **lignocellulosic biomass**."

Topic 2



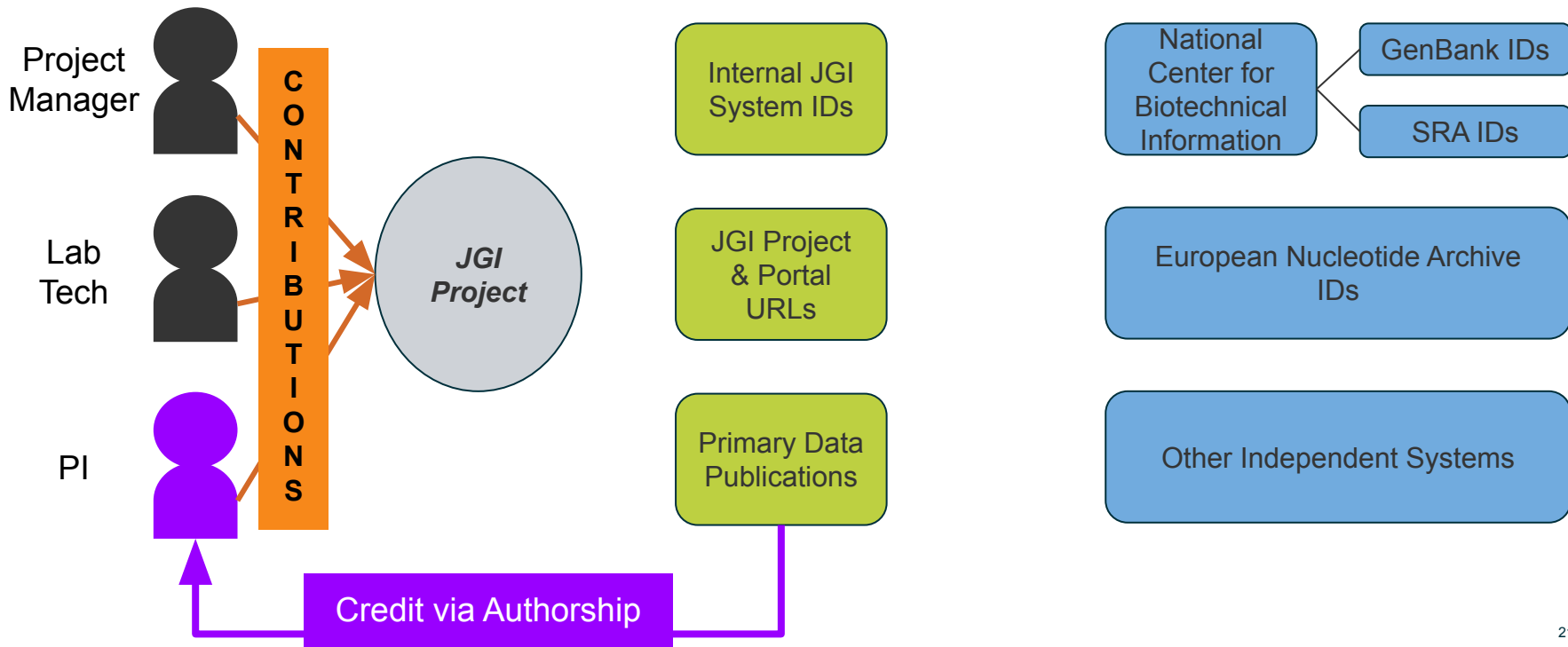
- 137 (25%) Publications
- 2.3k citations
- Top Journals: *Biotechnology for Biofuels*; *Bioresource Technology*

Citing Literature Share per Topic



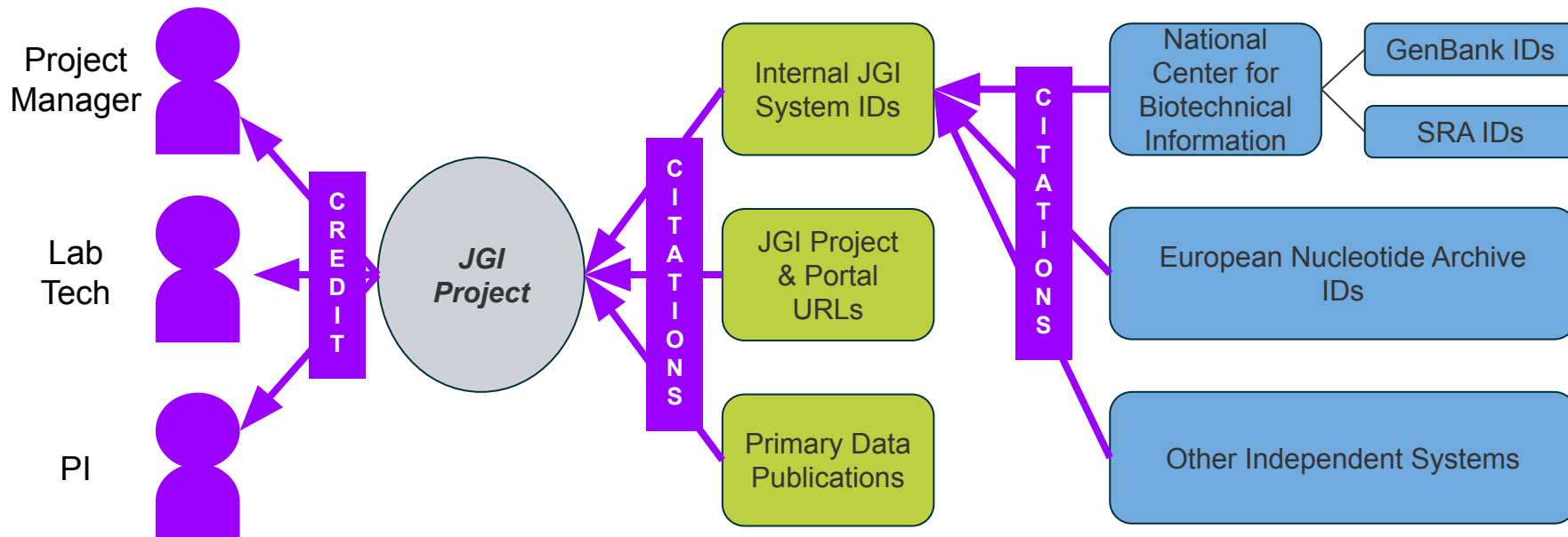
Equitable Contributor Metrics

Authorship Model - Often arbitrary and uneven credit attribution



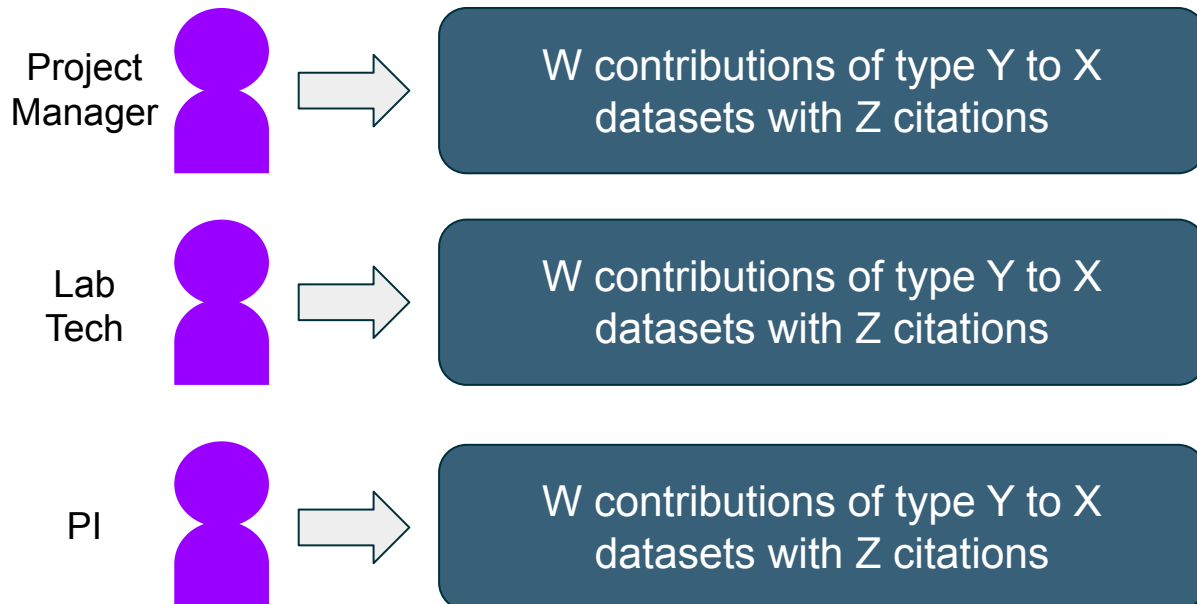
Equitable Contributor Metrics

Data Citation Model - Equitable attribution of credit



Equitable Contributor Metrics

Data Citation Model - Equitable attribution of credit



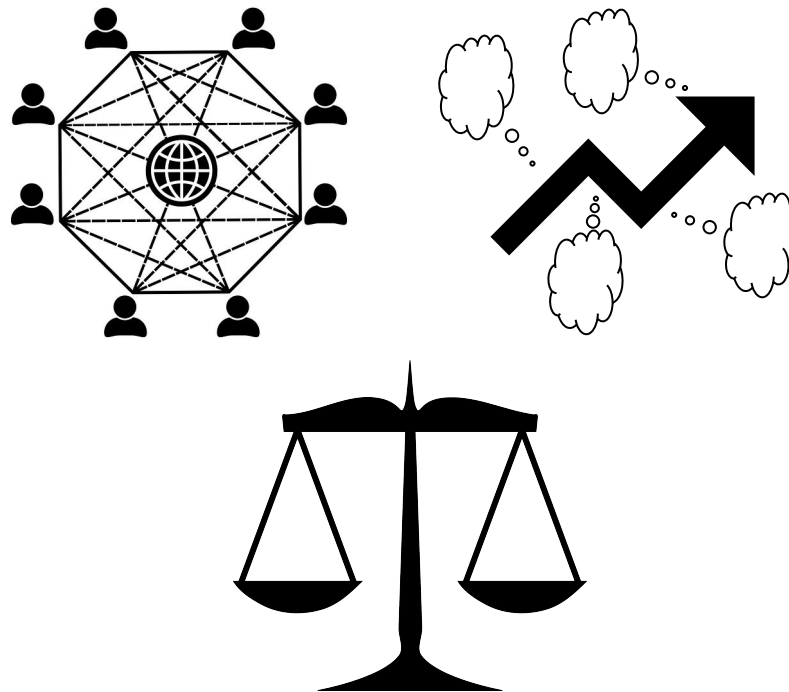
Example:

Hood & Sutherland,

"The data-index: An author-level metric that values impactful data and incentivizes data sharing"

Ecology and Evolution
Oct. 2021

- **GCS Goals**
 - Link metadata and capture data citations at scale
- **Results**
 - Many JGI data citations identified with high validity rates
- **Impact Implications**
 - More comprehensive picture of JGI community impact
 - Equitable contributor metrics



Thank you!

Genome Citation Service Team:



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TBK Reddy
Neil Byers



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Charles
Parker



George
Garrity



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