

# The AgBioData Consortium and Research Coordination Network:

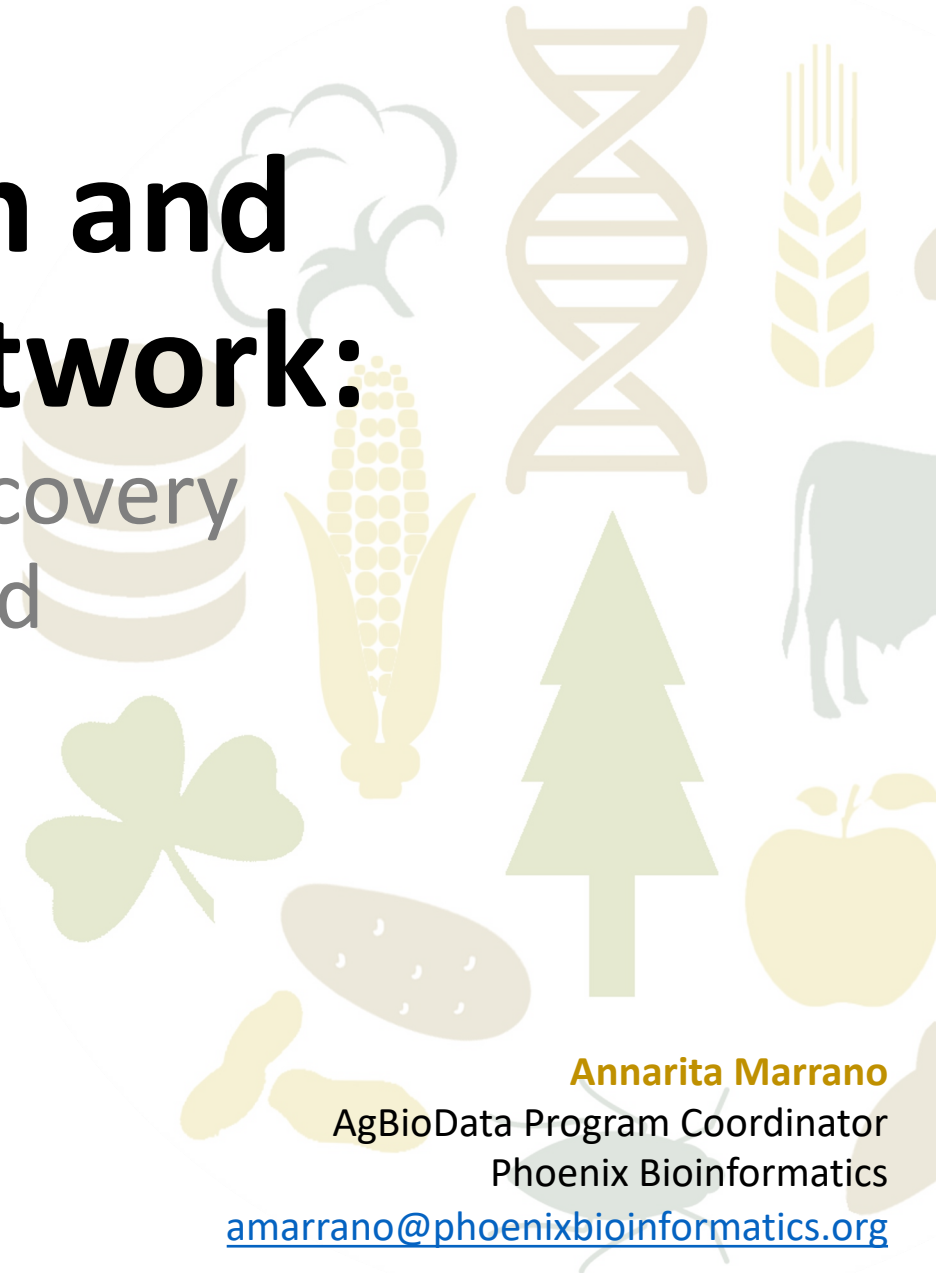
Enhancing Agricultural Research and Discovery  
through Improved Data Management and  
Education

*Plant Bioinformatics Resources for FAIR Agricultural Data Discovery and  
Reuse Workshop*

*Plant Biology 2023 Conference*

*Savannah (GA)  
August 6, 2023*

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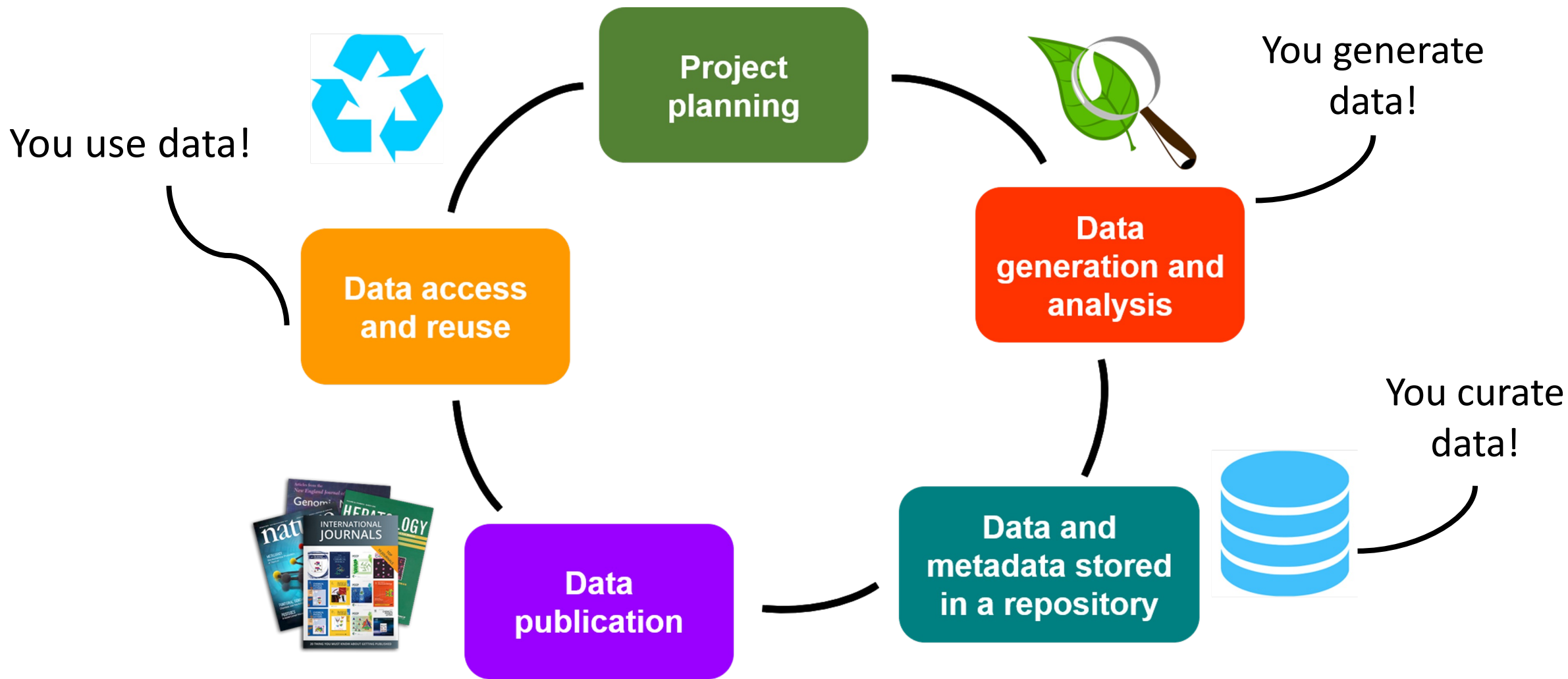


# The AgBioData Consortium

- ★ Founded in 2015
- ★ 45 Genetic, Genomics, and Breeding (GGB) resources
- ★ Over 250 members
- ★ **Mission:** *ensure standards and best practices for the acquisition, display, and retrieval of genomic, genetic, and breeding data.*



# How do the FAIR principle relates to you?



## What can you do to make your data FAIR?

- ❑ Become familiar and understand the FAIR principles
- ❑ Submit your data to a stable, long-term repository
- ❑ Include complete and detailed metadata with your data

## How AgBioData can help you?

- ✓ Webinars and a network of experts in the field for support
- ✓ Education Curriculum on databases and FAIR data
- ✓ Our member databases hosts different types of data and provide guidelines and support on what, how, and where to share

## What can you do to make your data FAIR?

- If there are community-standards, apply them when formatting your data
- Use established nomenclature guidelines for your species if they exist
- Release your data with clearly defined terms of use

## How AgBioData can help you?

- ✓ The member databases are working with the different stakeholders to establish community-standards in agricultural research
- ✓ Contact AgBioData/Community DB curators if you have any questions or seek for support with data management (e.g., DMP for grant proposals, etc.)

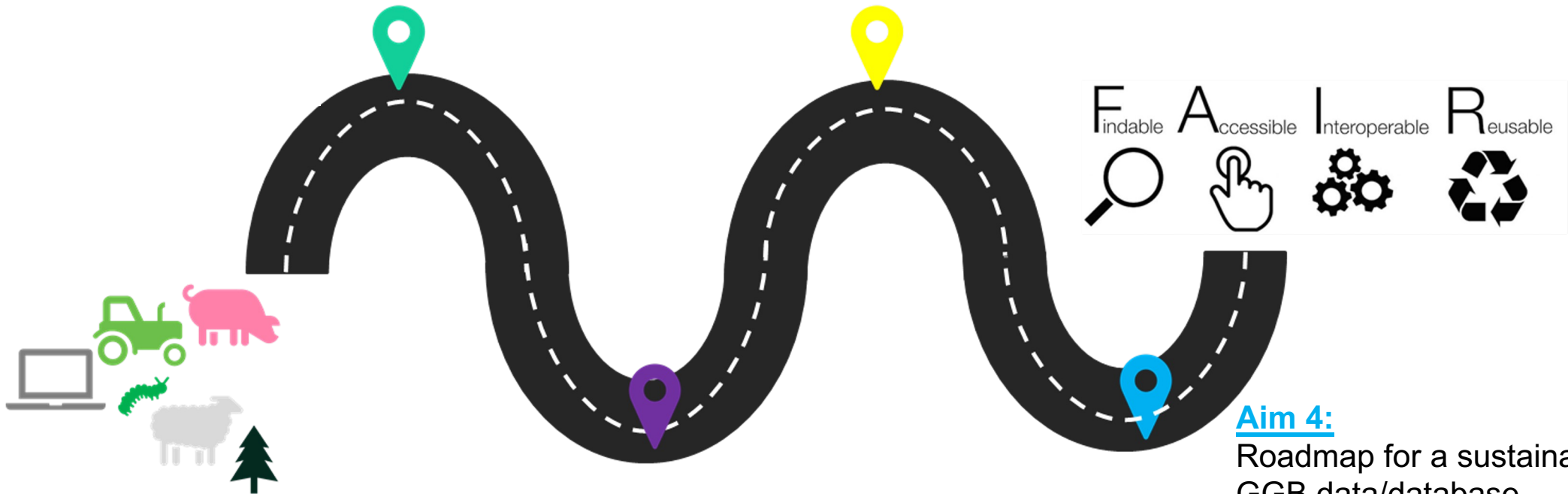


**Aim 1:**

Recommendations, standards, and implementation plans for FAIR data.

**Aim 3:**

Educational and training materials for researchers.



**Aim 2:**

Expand the network to include key stakeholders.

**Aim 4:**

Roadmap for a sustainable GGB data/database ecosystem.

# Past working groups & deliverables

- **Genotype to Phenotype**

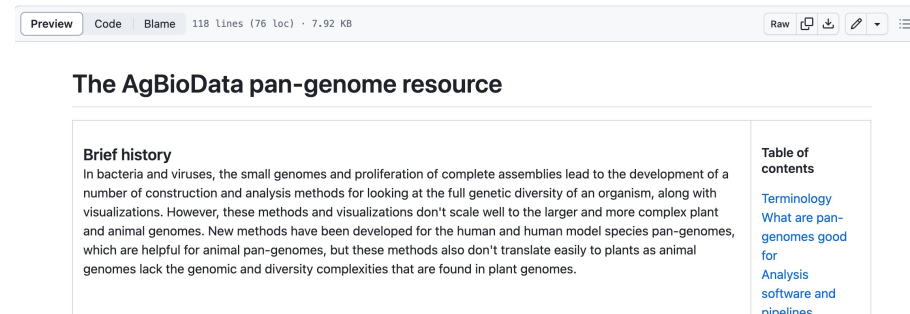
*Review*

## **Agricultural sciences in the big data era: Genotype and Phenotype Data Standardization, Utilization and Integration**

Cecilia H. Deng<sup>1\*</sup>, Sushma Naithani<sup>2\*</sup>, Sunita Kumari<sup>3\*</sup>, Irene Cobo-Simón<sup>4</sup>, Elsa H. Quezada-Rodríguez<sup>5,6</sup>, Maria Skrabisova<sup>7</sup>, Nick Gladman<sup>3,8</sup>, Melanie J. Correll<sup>9</sup>, Akeem Babatunde Sikiru<sup>10</sup>, Olusola O Afuwape<sup>11</sup>, Annarita Marrano<sup>12</sup>, Ines Rebollo<sup>13</sup>, Wentao Zhang<sup>14</sup> and Sook Jung<sup>15\*</sup>

*Preprints;* <https://doi.org/10.20944/preprints202306.1013.v1>

- **Pan-genomes**



The screenshot shows a GitHub repository page for 'The AgBioData pan-genome resource'. The repository has 118 lines of code (76 LOC) and is 7.92 KB in size. The page includes a 'Brief history' section, a 'Table of contents' section, and a 'What are pan-genomes good for' section. The 'Brief history' section states: 'In bacteria and viruses, the small genomes and proliferation of complete assemblies lead to the development of a number of construction and analysis methods for looking at the full genetic diversity of an organism, along with visualizations. However, these methods and visualizations don't scale well to the larger and more complex plant and animal genomes. New methods have been developed for the human and human model species pan-genomes, which are helpful for animal pan-genomes, but these methods also don't translate easily to plants as animal genomes lack the genomic and diversity complexities that are found in plant genomes.' The 'Table of contents' section includes links to 'Terminology', 'What are pan-genomes good for', and 'Analysis software and pipelines'.

[https://github.com/AgBioData/Pan-genomes/blob/main/Pan-genome\\_resource.md](https://github.com/AgBioData/Pan-genomes/blob/main/Pan-genome_resource.md)

- **Data Federation**
- **Ontologies**

*[Submitted on 18 Jul 2023]*

## **Data sharing and ontology use among agricultural genetics, genomics, and breeding databases and resources of the AgBioData Consortium**

Jennifer L. Clarke, Lauree D. Cooper, Monica F. Poelchau, Tanya Z. Berardini, Justin Elser, Andrew D. Farmer, Stephen Ficklin, Sunita Kumari, Marie-Angélique Laporte, Rex T. Nelson, Rie Sadohara, Peter Selby, Anne E. Thessen, Brandon Whitehead, Taner Z. Sen

Over the last several decades, there has been rapid growth in the number and scope of agricultural genetics, genomics and breeding (GGB) databases and resources. The AgBioData Consortium ([this https URL](https://www.agbiodata.com/)) currently represents 44 databases and resources covering model or crop plant and animal GGB data, ontologies, pathways, genetic variation and breeding platforms (referred to as 'databases' throughout). One of the goals of the Consortium is to facilitate FAIR (Findable, Accessible, Interoperable, and

<https://doi.org/10.48550/arXiv.2307.08958>

# Our current working groups

New working groups coming soon!

- Data Federation Training
- Data Reuse
- FAIR Scientific Literature
- Genome Assembly and Annotation Nomenclature
- Public Genetic Resources
- Standards for genetic variation

**Aim 1**

- Diversity, Equity, and Inclusion (DEI) Recruiting

**Aim 2**

- Education

Education Curriculum on databases and FAIR data

**Aim 3**

- Sustainability

**Aim 4**



# Community workshops and outreach

*Bring together the working groups and all the stakeholders that generate, use, curate, and manage data in agriculture*

- 2022 AgBioData Community Workshop (March 15-17; Virtual)
- The AgBioData Consortium: Challenges and Recommendations for FAIR Genetic, Genomic and Breeding Data” at PAG 30 (San Diego, January 2023)
- 2023 AgBioData Community Workshop (May 1-2; Chicago – Hybrid)
- “Bringing FAIR Data in the Classroom” at Plant Biology 2023 Conference (August 5; Virtual)



**2024 Community Workshop  
Virtual – Spring 2024!!!!**

# How to participate in AgBioData:

- **Interested in joining our working groups?**

**Do you need help with your data?**

Send an email to [agbiodata@gmail.com](mailto:agbiodata@gmail.com)!



- **Become a member!**

Visit our website [www.agbiodata.org](http://www.agbiodata.org)

- Join our Slack workplace!



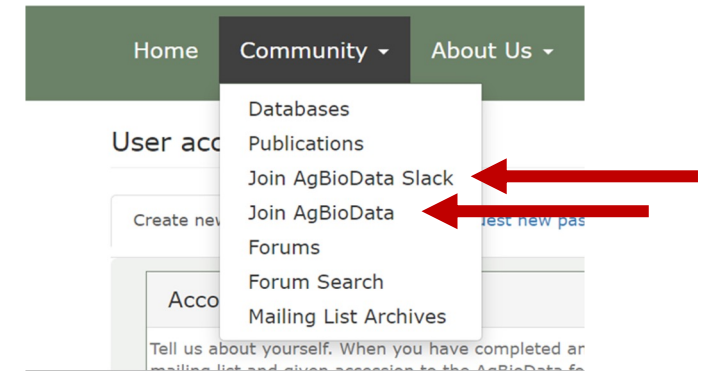
- @AgBioData 

- Follow us on LinkedIn 

- Monthly meetings/webinars (1<sup>st</sup> Wed of the month)



- **If you have a GGB resource, join the consortium!**



Everything you need in one place!

<https://linktr.ee/agbiodata>

**Poster  
# 500-22**

**Booth  
# 406**

# Acknowledgements

## AgBioData SC members:

Jacqueline Campbell  
Peter Harrison  
Sunita Kumari  
Dorrie Main  
John P. McNamara  
Sushma Naithani  
Monica Poelchau  
Leonore Reiser  
Meg Staton

**The AgBioData  
consortium**

## Past AgBioData SC members:

Ethy Cannon  
Laurel Cooper  
Lisa Harper  
Eva Huala  
Sook Jung  
Marcela Tello-Ruiz

## Past PC:

Darwin Campbell

