



जैवप्रौद्योगिकी विभाग
DEPARTMENT OF
BIOTECHNOLOGY

सत्यमेव जयते



United Nations
Educational, Scientific and
Cultural Organization



क्षेत्रीय जैव प्रौद्योगिकी केन्द्र
Regional Centre
for Biotechnology

INDIAN CROP PHENOME DATABASE

Advancing Crop Research Through Open Phenomic Data



Presented by: Dr. Sonia Balyan
On behalf of Team ICPD



CONTACT US



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Website:

<https://ibdc.dbtindia.gov.in/icpd/>





AgBioData

Toward enhanced genomics, genetics, and breeding research outcomes through standardization of practices and protocols across agricultural databases





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International Centre for Genetic
Engineering and Biotechnology

NATIONAL
INFORMATICS
CENTRE



National Institute of Immunology



United Nations
Educational, Scientific and
Cultural Organization



क्षेत्रीय जैव प्रौद्योगिकी केन्द्र
Regional Centre
for Biotechnology



Indian Biological Data Centre

Regional Centre for Biotechnology,
NCR Biotech Science Cluster, 3rd Milestone,
Faridabad-Gurugram Expressway,
Faridabad – 121 001



<https://ibdc.dbtindia.gov.in/>



support@ibdc.rcb.res.in



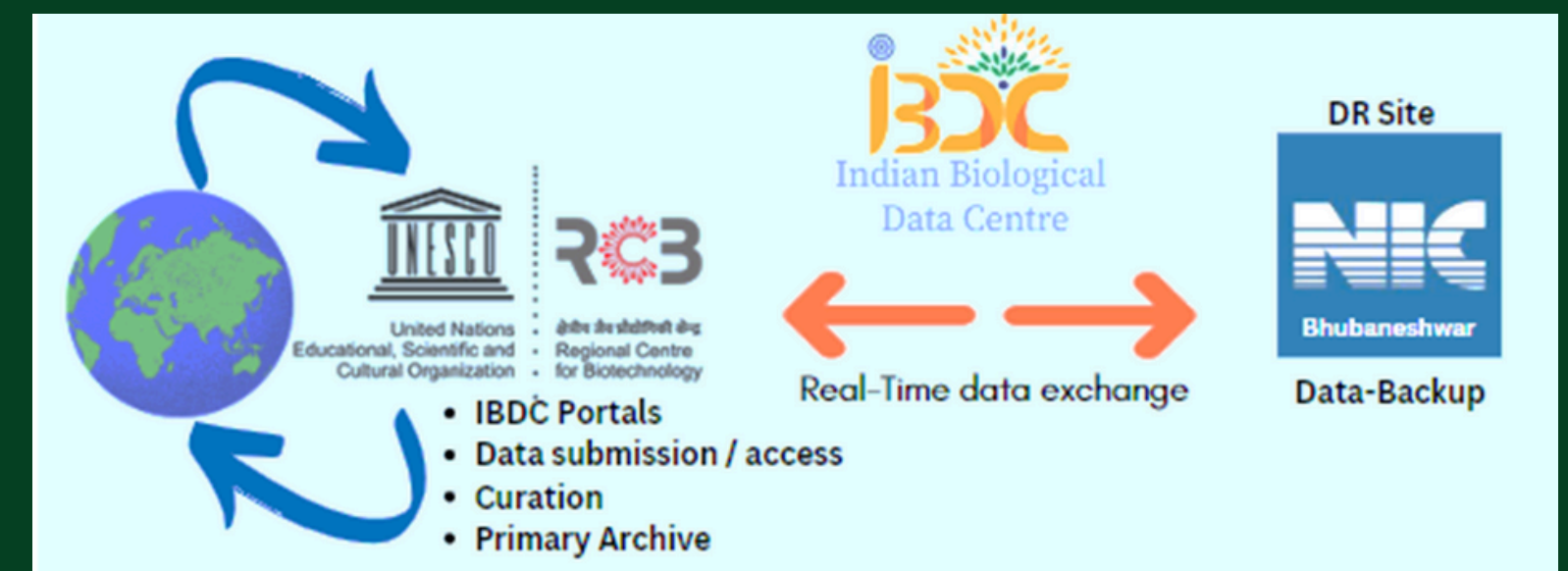
@ibdctweets





Indian Biological Data Centre

- IBDC is the first National life science data repository in India.
- It is supported by the **Government of India (GOI)** through the **Department of Biotechnology (DBT)**.
- It enables the implementation of the “**Biotech-Pride Guidelines**”.
- It is a joint project of **Regional Centre for Biotechnology, National Institute of Immunology, International Centre for Genetic Engineering & Biotechnology** and **National Informatics Centre**.



LOCATION

The IBDC was dedicated to the nation by **Dr. Jitendra Singh**, Honorable Minister for Science and Technology, GOI on Nov 10, 2022.



Indian Biological Data Center,
Regional Centre for Biotechnology,
NCR Biotech Science Cluster, 3rd Milestone,
Faridabad-Gurugram Expressway,
Faridabad – 121 001



<https://ibdc.dbtindia.gov.in/>



support@ibdc.rcb.res.in

It is housed at Regional Centre of Biotechnology, an academic institution established by the Department of Biotechnology, Govt. of India with regional and global partnerships synergizing with the programmes of UNESCO as a Category II Centre.

DEDICATION

The IBDC was dedicated to the nation by **Dr. Jitendra Singh**, Honorable Minister for Science and Technology, GOI on Nov 10, 2022.

Business Standard

Saturday, December 3, 2022 | 13:54 IST English | Hindi

Centre unveils India's first national repository for life science data

Data submission and access portals for other data types are under development and would be launched shortly

Data

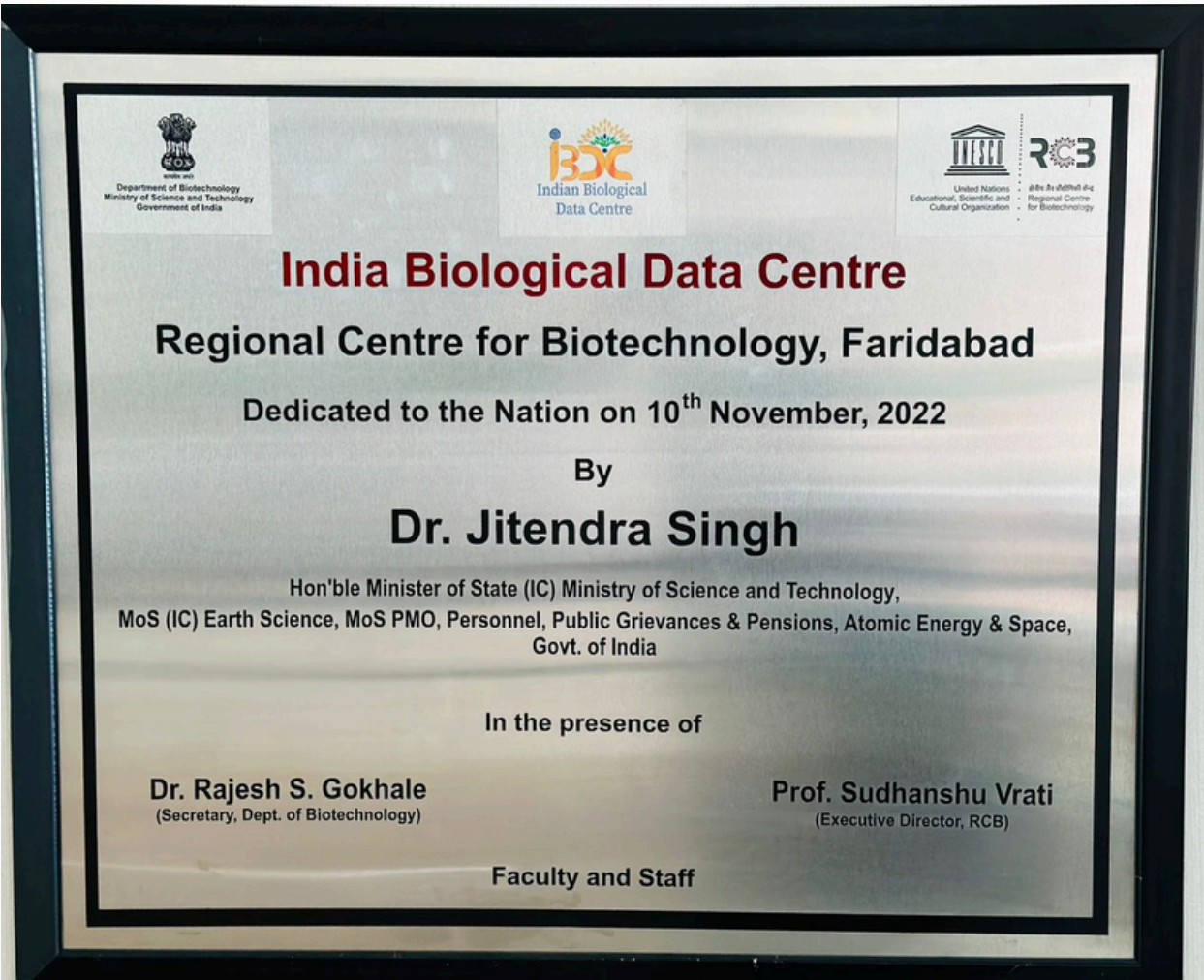
The 'Indian Biological Data Centre' is established at Regional Centre of Biotechnology in Faridabad

India News | Press Trust of India | Updated: November 13, 2022 11:17 am IST



IBDC: Government launches India's first national repository for life science data

As per the BIOTECH-PRIDE guidelines of the Government of India, IBDC is mandated to archive all life science data generated from publicly-funded research in India.



MISSION



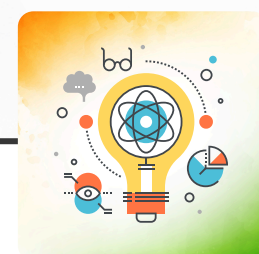
WHAT WE DO?



**Biological Data
Archive
Resources**



**High
Performance
Computational
Facility**



**Support to
Project of
National
Importance**



**Big Data
Analytics**



User Resources



**Trainings:
Data
Submission &
Data Analysis**



Projects of National Importance



260,448 submissions from 60 institutes



9768 samples with (gvcf, phenotype and raw data of ~900TB of data size)



9833 samples submitted



557 samples submitted

Major Mission Projects

Triticum aestivum

Germplasm Characterization and Trait Discovery in Wheat using Genomics Approaches and its Integration for Improving Climate Resilience, Productivity and Nutritional quality.

Oryza sativa

Mainstreaming rice landraces diversity in varietal development through genome-wide association studies: A model for large-scale utilization of gene bank collections of rice.

Linum usitatissimum

Leveraging genetic resources for accelerated genetic improvement of Linseed using comprehensive genomics and phenotyping approaches.

Cicer arietinum

Characterization of Chickpea Germplasm Resource to Accelerate Genomics-assisted Crop Improvement.

Carthamus tinctorius

Exploiting Genetic Diversity for Improvement of Safflower through Genomics Assisted Discovery of QTLs/Genes Associated with Agronomic Traits.

Sesamum indicum

Mainstreaming sesame germplasm for productivity enhancement and sustainability through genomics assisted core development and trait discovery.

IBDC Portals



Indian Biological Images Archive

- An online repository of biological images generated from multiple imaging modalities, e.g., histopathology, X-ray, etc.
- Total submissions: 241020 Total data size: 557.25 GB
- No. of registered users: 67 from 47 organizations.
- <https://ibdc.dbtindia.gov.in/ibia/>



Indian Metabolome Data Archive

- Catalogue metabolomics data and associated metadata generated through the analytical techniques MS and NMR.
- Total submissions: 1264 Total data size: 10.6 GB
- No. of registered users: 56 from 51 organizations.
- <https://ibdc.dbtindia.gov.in/imda/>



Indian Crop Phenome Database

- Central portal for the archival and access of crop phenotype data.
- Total submissions: 367 Total data size: 317MB
- No. of registered users: 74 from 42 organizations.
- <https://ibdc.dbtindia.gov.in/icpd/>



Indian Nucleotide Data Archive

- The Indian Nucleotide Data Archive (INDA), is an open-access platform for archival of nucleotide sequencing data which provides Internationally recognized INSDC and IBDC accessions.
- Total submissions: 23099 Total data size: 44.7 TB
- No. of registered users: 325 from 105 organizations.
- <https://ibdc.dbtindia.gov.in/inda/>



Indian Proteome Databank

- IPD is a MS-Proteomics data submission and archival portal.
- Total submissions: 118 Total data size: 6200GB
- No. of registered users: 58 from 41 organizations.
- <https://ibdc.dbtindia.gov.in/ipd/>



Indian Structural Data Archive

- ISDA is an archive of three-dimensional structures of biological macromolecules with function annotations from various sources.
- Total submissions: 239492 Total data size: 30000GB
- No. of Indian entries: 7145 from 65 organizations.
- <https://ibdc.dbtindia.gov.in/isda/>



INDA-Controlled Access

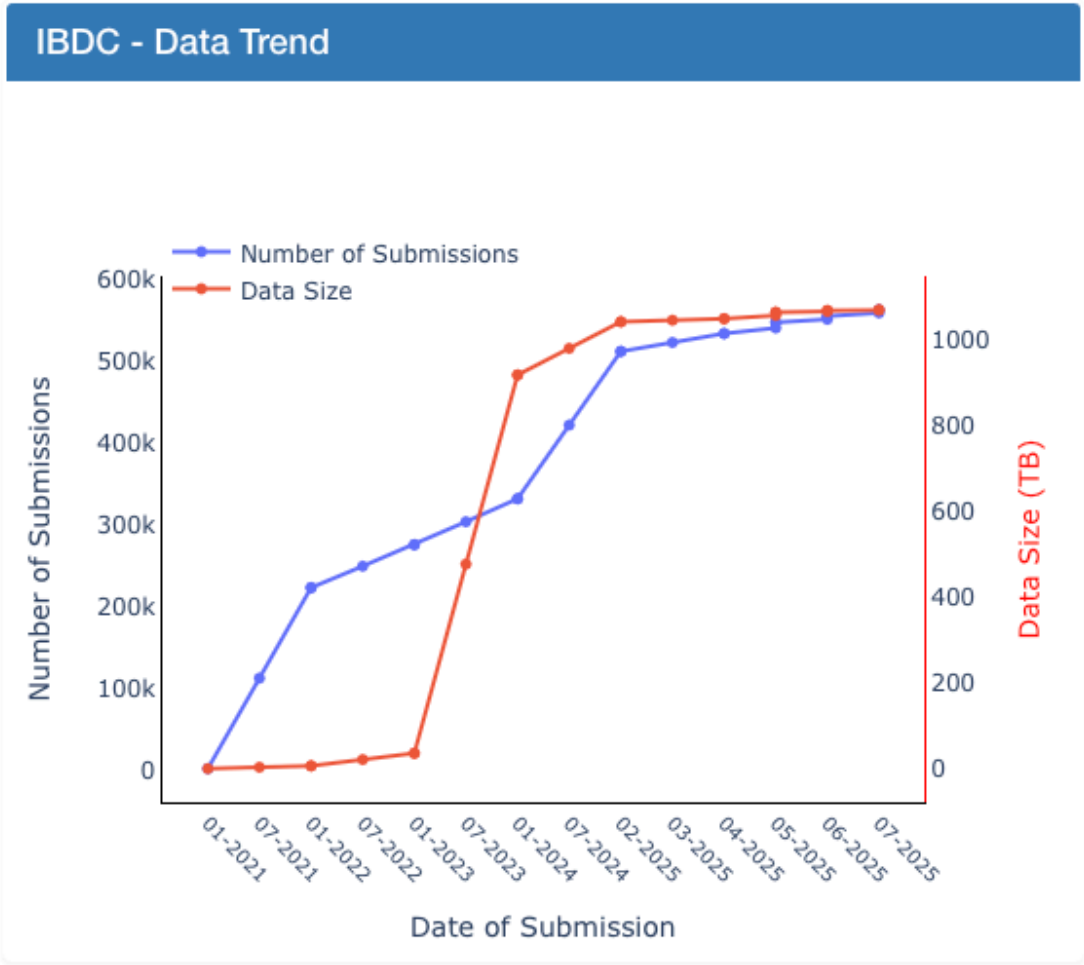
- The Indian Nucleotide Data Archive- Controlled access (INDA-CA), is a controlled access platform for archiving nucleotide sequencing data.
- Total submissions: 295666 Total data size: 1012 TB
- No. of registered users: 414 from 200 organizations.
- <https://ibdc.dbtindia.gov.in/indasecure/>



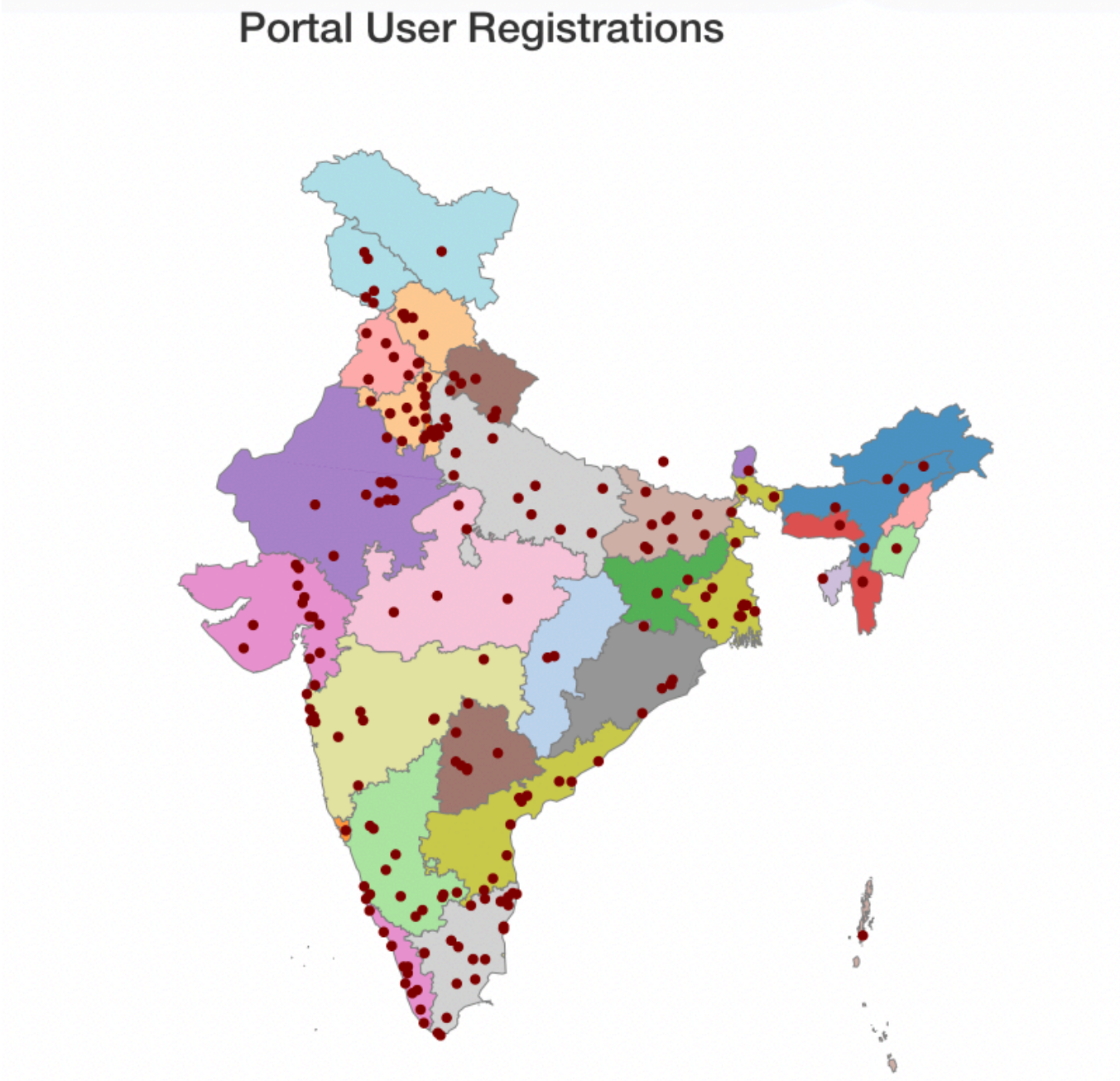
Trainings



Conferences



Overall Data	Value
No. of Users	1003
No. of Submission	576607
Total Downloads	23746684
Total Visitors	448107
No. of Open Access	265400
No. of Restricted Access	9833
No. of Managed Access	301374



Data Portal Diversity @IBDC



ICPD



IAPD
Indian Animal Phenome Database



IHPD
Indian Human Phenome Database

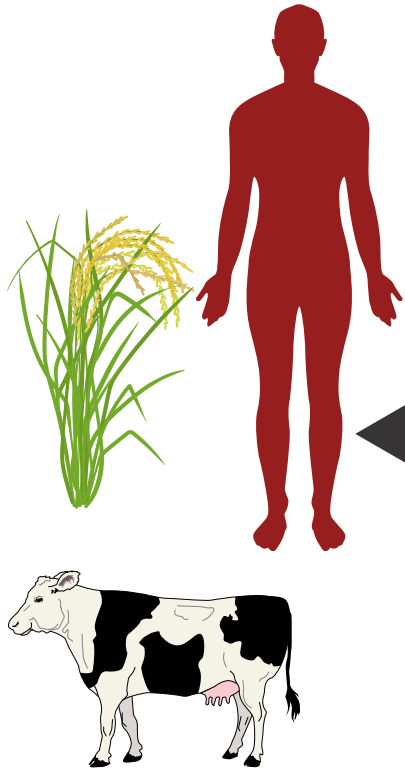


IMPD
Indian Microorganisms Phenome Database



IGR
Indian GWAS Repository

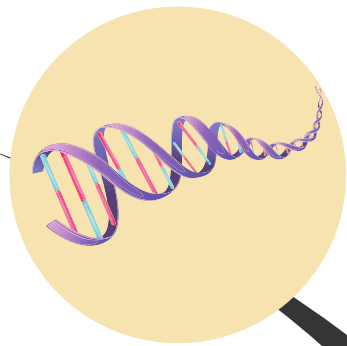
Phenotype




Images




IBIA
Indian Biological Images Archive




Genes, Genomes and Variation




INDA
Indian Nucleotide Data Archive



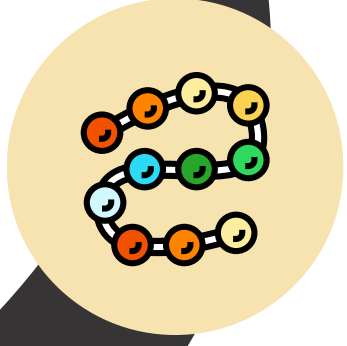
INDA-CA
Indian Nucleotide Data Archive
Controlled Access




Functional Genomics



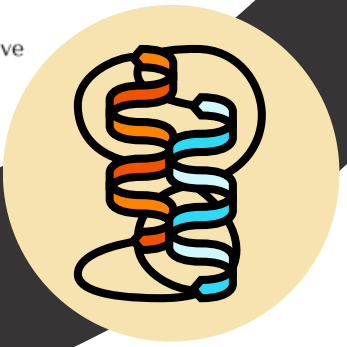
IADA
INDIAN ARRAY DATA ARCHIVE




Proteomics



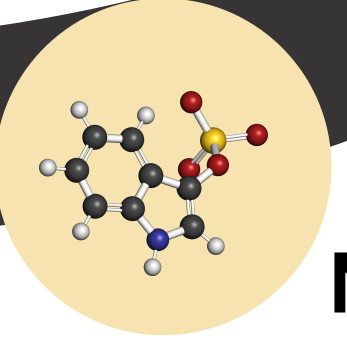
IPD
Indian Proteome Databank




Molecular Structures



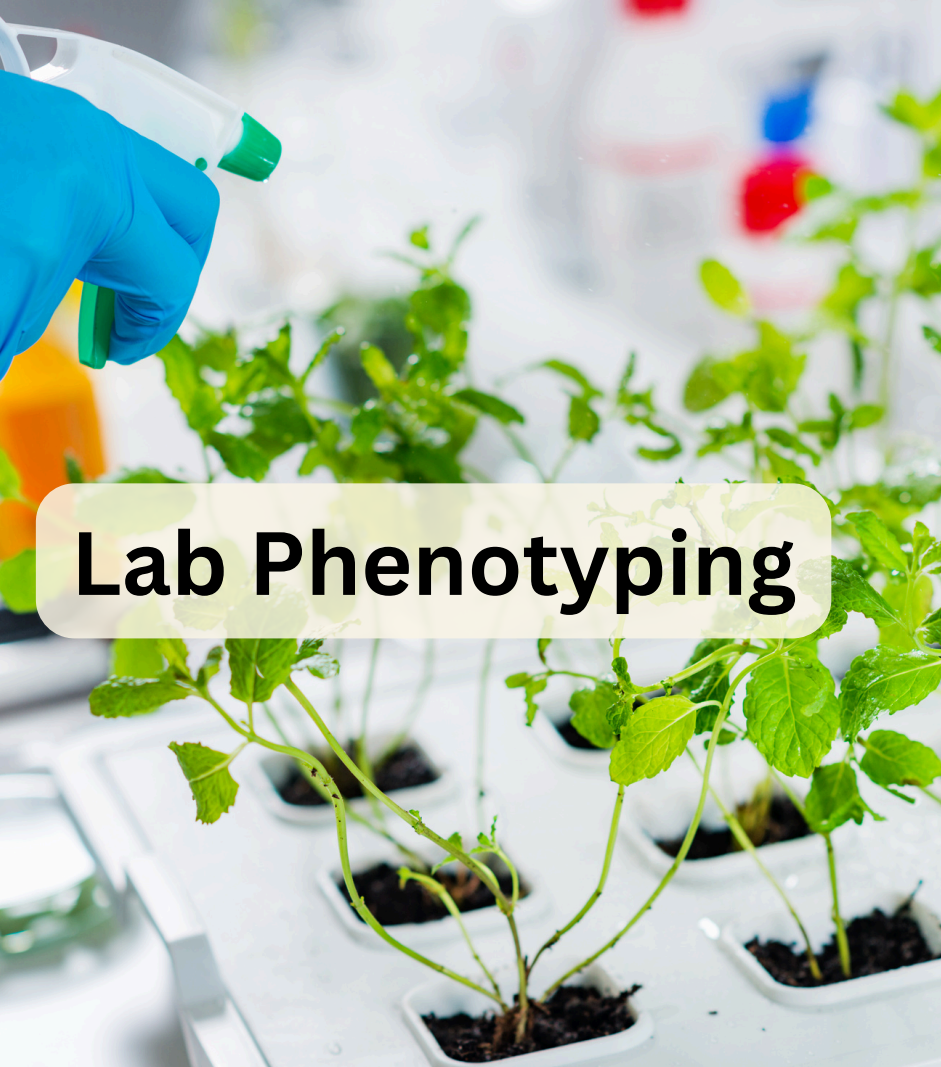
ISDA
Indian Structural Data Archive



Metabolomics



IMDA
Indian Metabolome Data Archive



Lab Phenotyping



**Field Phenotyping
(Semi-controlled)**



Field Phenotyping



**Phenotyping facility
(controlled environment)**

Levels of Phenotyping

Heterogeneous data

- Indoor/outdoors
- Collaborative projects
- Manual to automated data collection
- High throughput phenotyping

Increasing with technology

- Devices
- Sensors
- Measurements



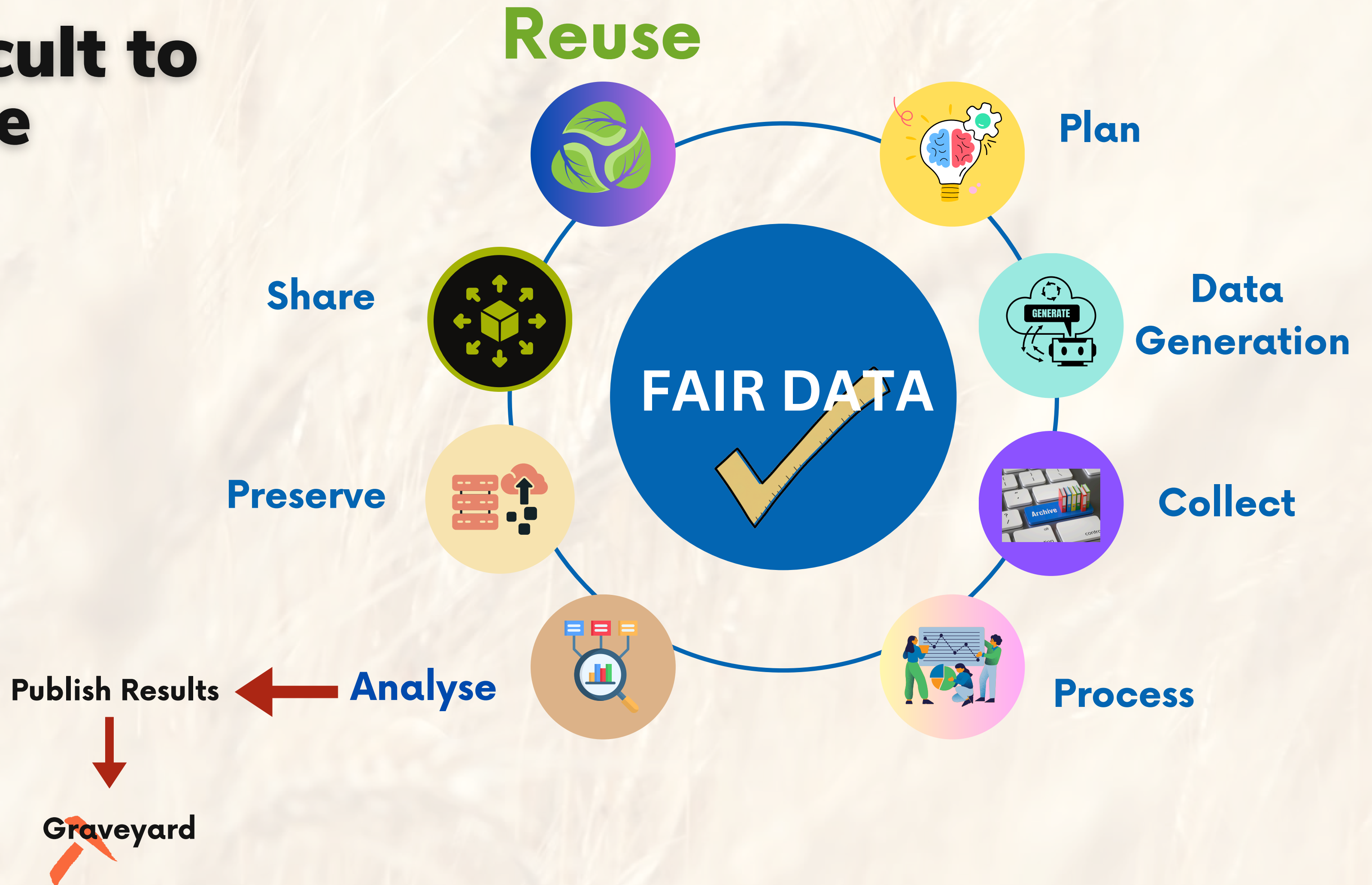
Complicated

- Different environments and ecosystems
- Different tissues and stages
- Different scales
- Different interactions
- varied names and measuring conventions

Poorly Documented

Lack of proper meta data formats and dedicated repositories

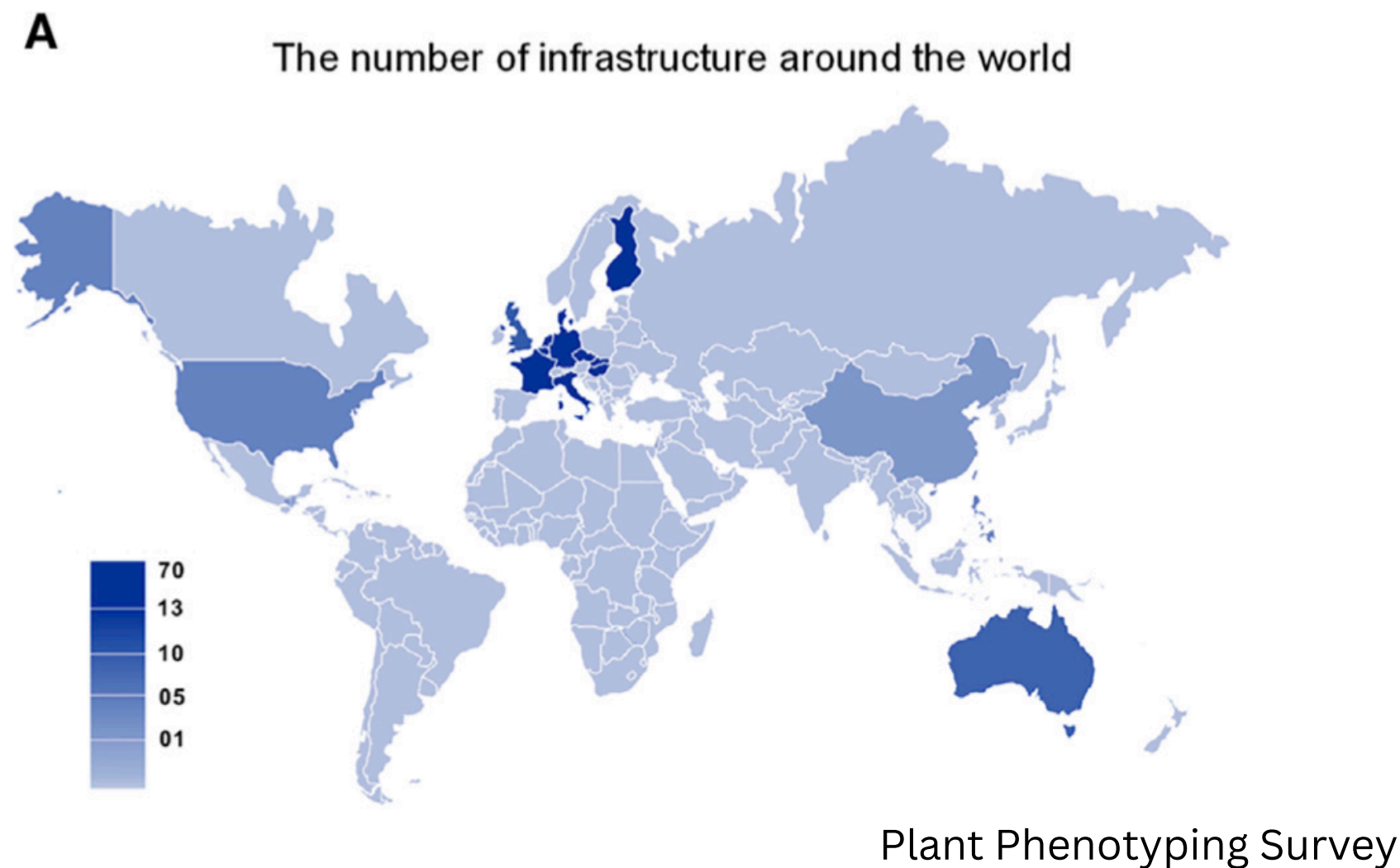
Difficult to Reuse



Fiorani and Schurr (2013) referred to plant phenotyping as the set of methodologies and protocols used to accurately measure plant growth, architecture, and composition at different scales.

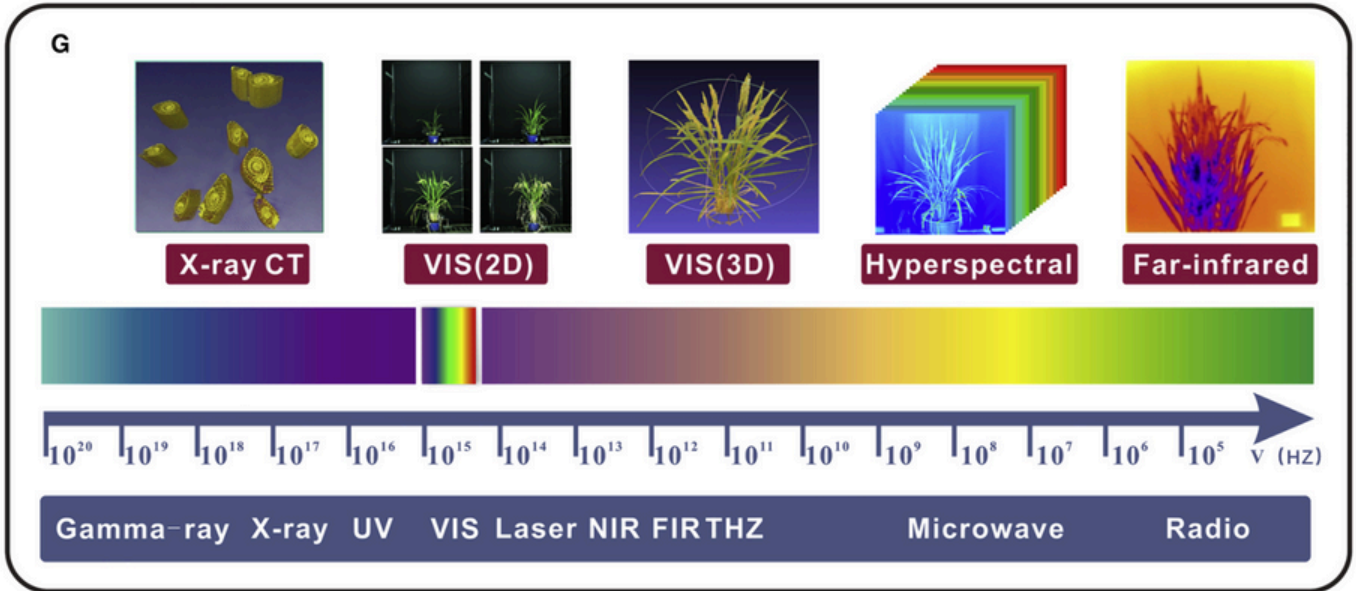
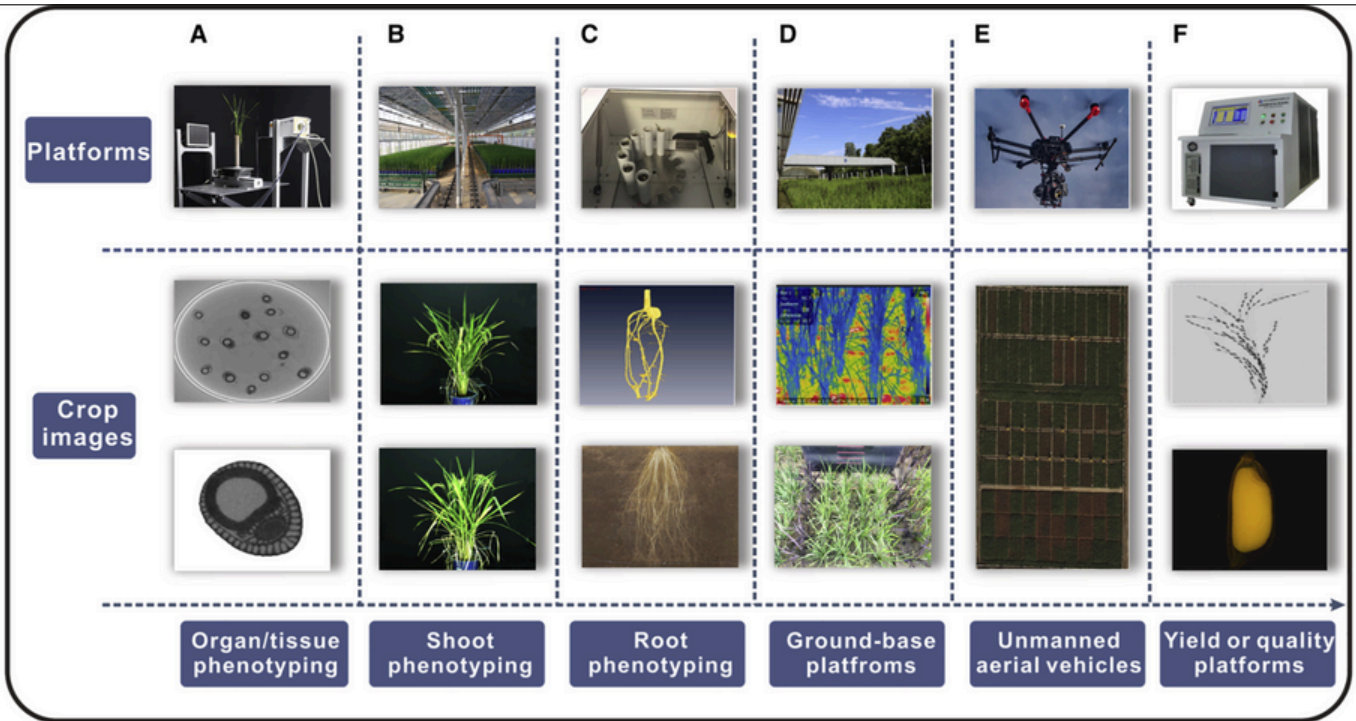
Phenotype Data
Generation

Traditional Phenotyping
High-Throughput Phenotyping (HTP)



High throughput phenotyping in India

- Nanaji Deshmukh Plant Phenomics Centre (NDPPC) at Indian Agricultural Research Institute (ICAR-IARI) in New Delhi,
- International Crops Research Institute for the Semi-Arid Tropics (ICRISAT)
- Indian Institute of Horticultural Research (IIHR): Based in Bengaluru, Karnataka



Data and Observations

Importance for India

1. Employment Contribution:

- **Workforce Engagement:** Approximately 45.5% of India's total workforce is employed in agriculture and related activities. Statista
- **Livelihood Support:** The agriculture sector provides livelihood support to about 42.3% of the population.

2. GDP Contribution:

- **Gross Value Added (GVA):** In the fiscal year 2022-23, agriculture contributed **18.3%** to India's GVA.
- **Overall GDP Share:** For the fiscal year 2024-25 (provisional estimates), agriculture's share in the country's GDP is approximately **16%**. India Budget

3. Food Security:

- **Undernourishment:** Despite achieving food sufficiency, India is home to over 190 million undernourished people, accounting for a quarter of the world's hungry population.

4. Growth Trends:

- **Sector Growth:** The agriculture sector has registered an average annual growth rate of 4.18% at constant prices over the last five years.

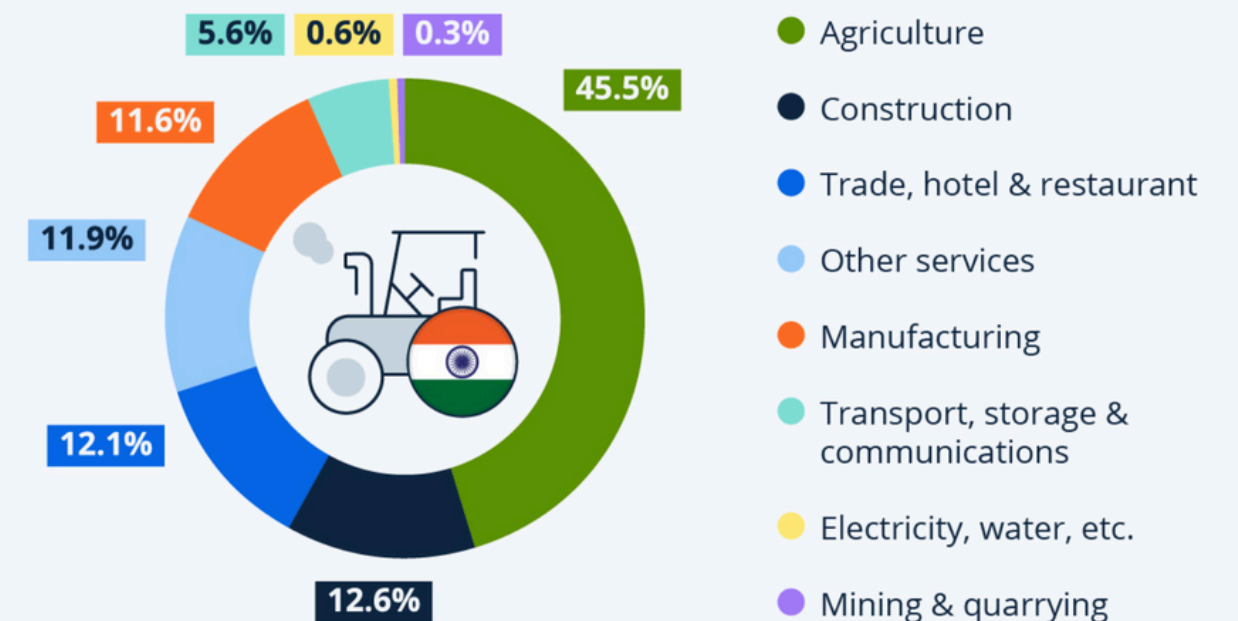
5. Budgetary Allocations:

- **Financial Year 2025-26:** The government has allocated approximately 1.75 trillion rupees (about \$20 billion) to agriculture sector

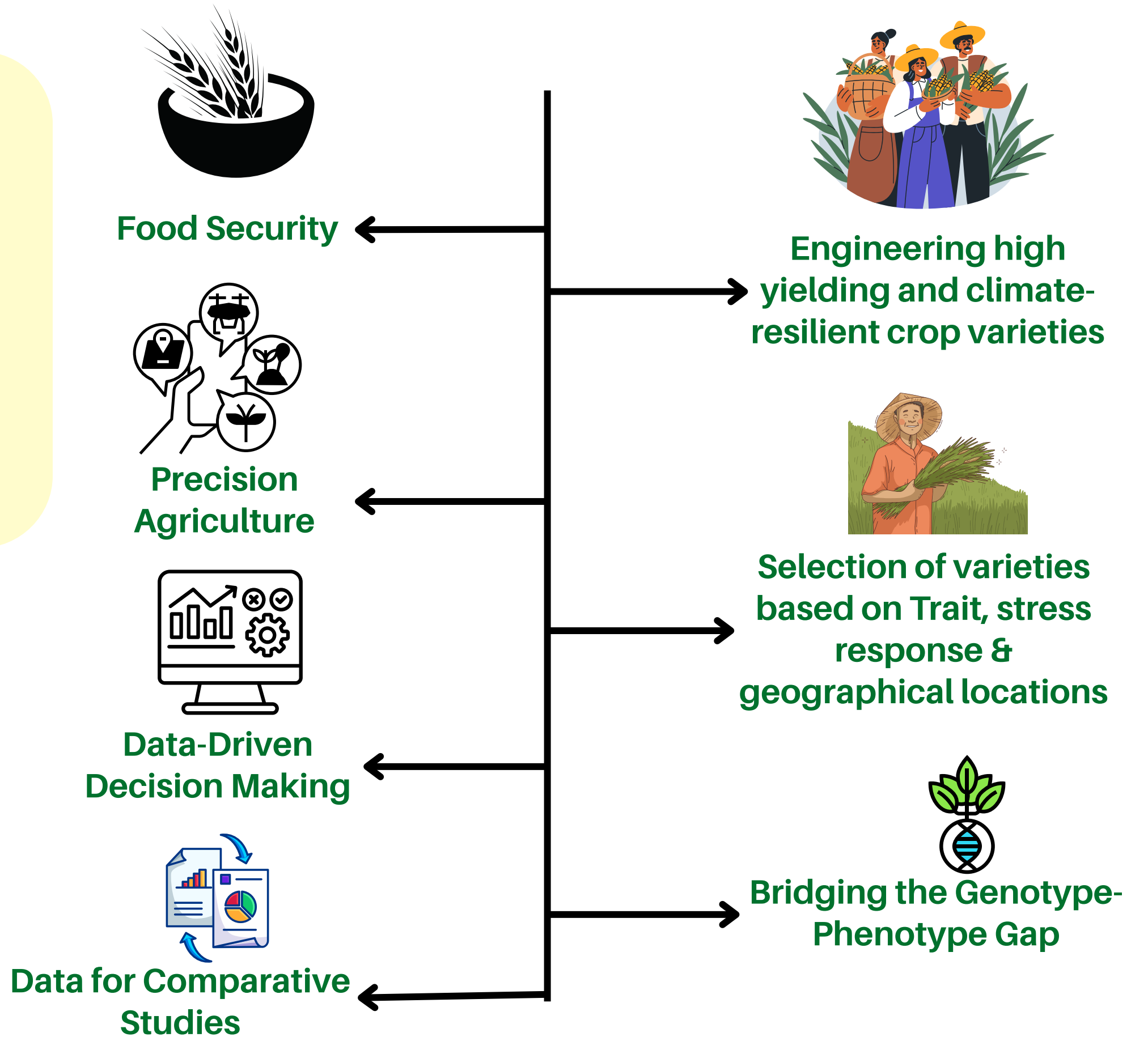
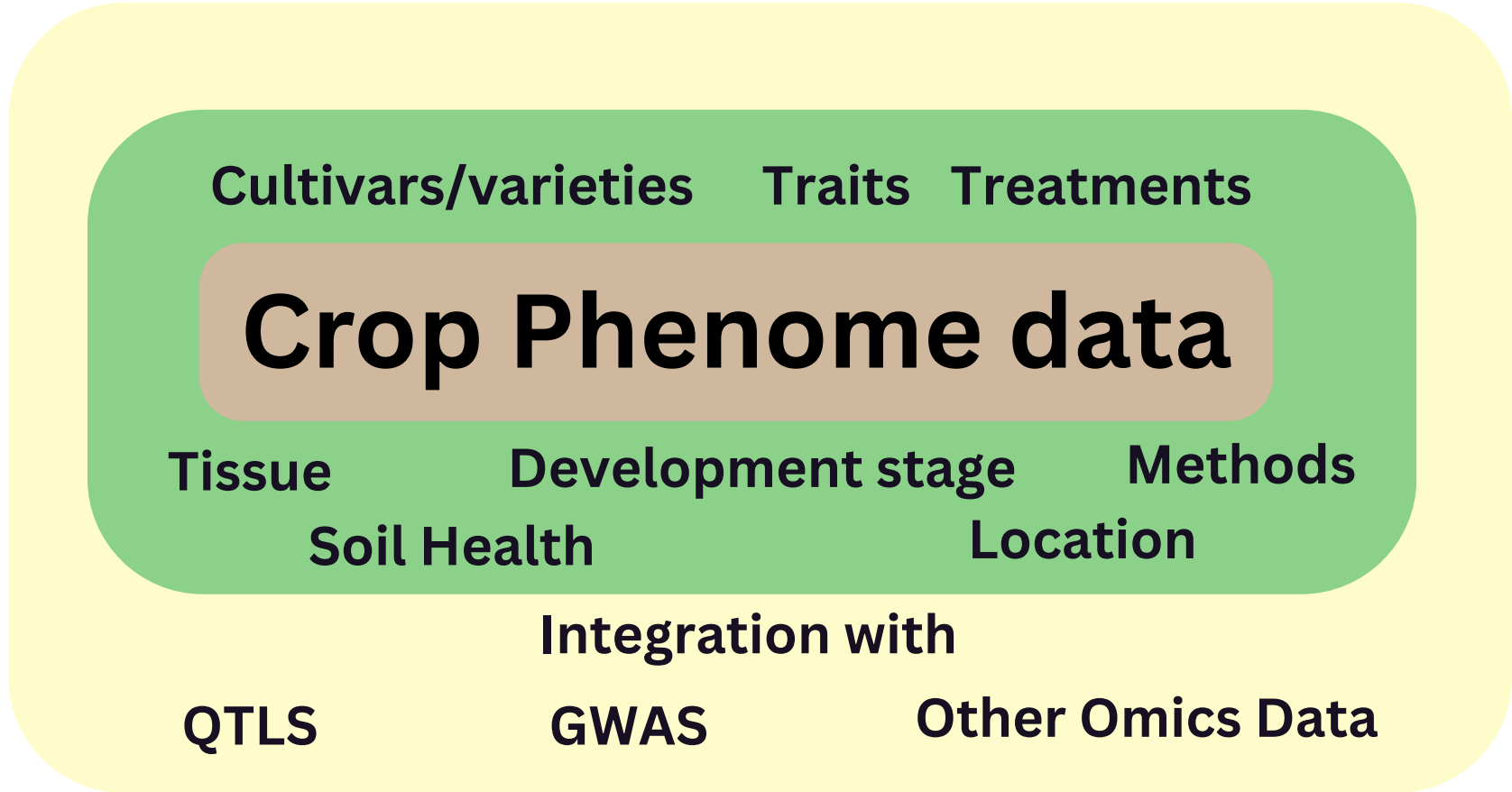


Agriculture Is the Biggest Employer in India

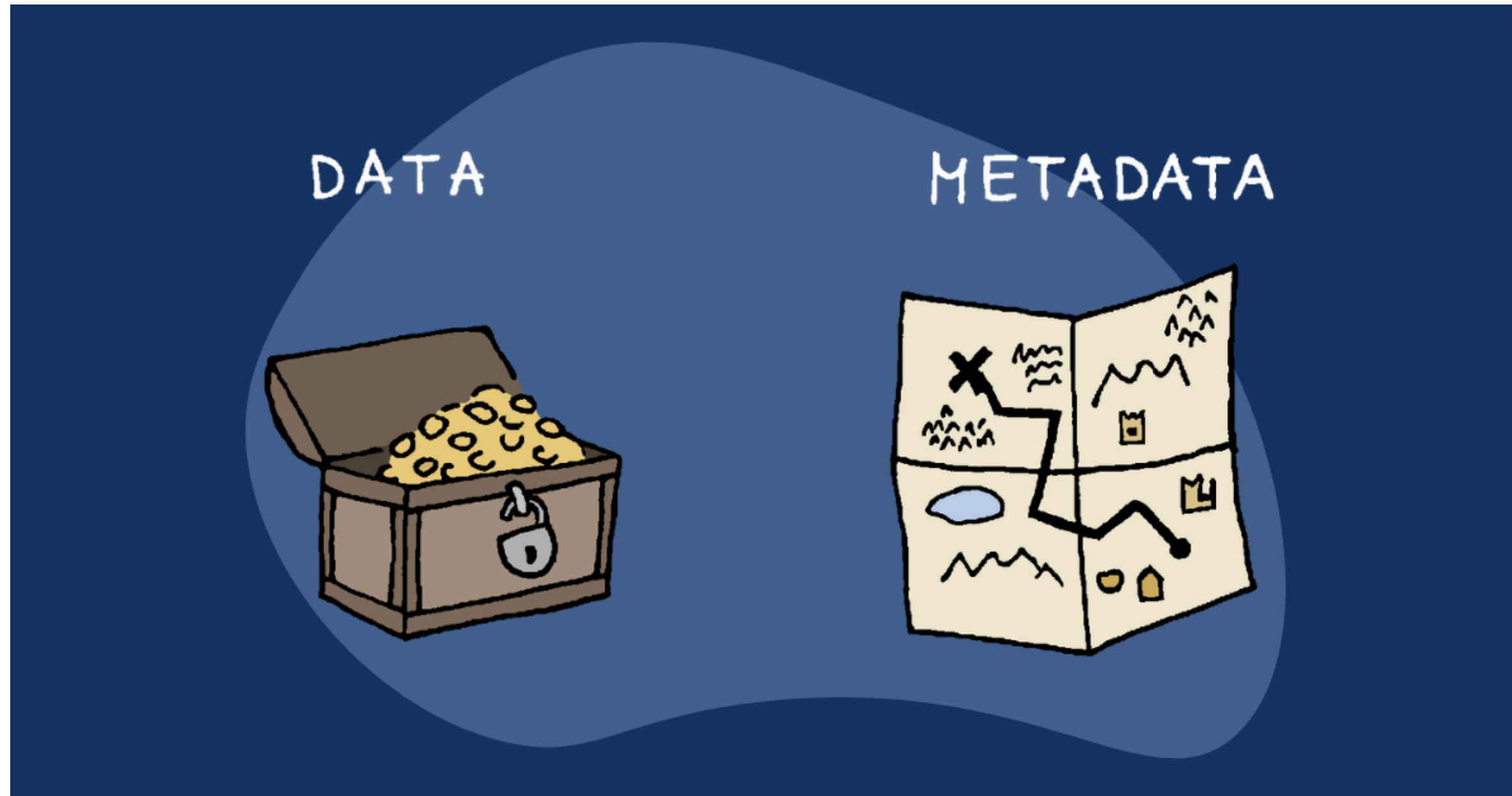
Share of workers in India, by broad industry of work (2021-22)



WHY PHENOTYPE DATA IS IMPORTANT



DATA ALONE IS NOT SUFFICIENT....



Meta-Data: Data about Data

Why Metadata Matters for Phenotype Data

🌱 Making sense of data

Metadata tells the story behind the numbers – what was measured, when, where, how, and why. It connects phenotypic traits with the actual experiment, environment, and methods used. Without metadata, phenotype data is incomplete and hard to trust.

Metadata Makes Data FAIR

- Findable – Clear descriptions help others discover your data easily.
- Accessible – Metadata explains how and where the data can be reached.
- Interoperable – Using shared standards and vocabularies lets data from different studies “talk” to each other.
- Reusable – With enough detail, your data can be reliably used, compared, and built upon in the future.

Why it's important

- ✓ Others can reproduce your work and validate findings.
- ✓ Studies from different sites and seasons can be compared fairly.
- ✓ Phenotype data can be integrated with genomic and environmental data.
- ✓ Good metadata turns raw data into long-lasting knowledge.



Indian Crop Phenome Database (ICPD)

We empower the researcher with various digitization services for harnessing their crop phenotyping projects.

[Submit Data](#)[Browse Data](#)

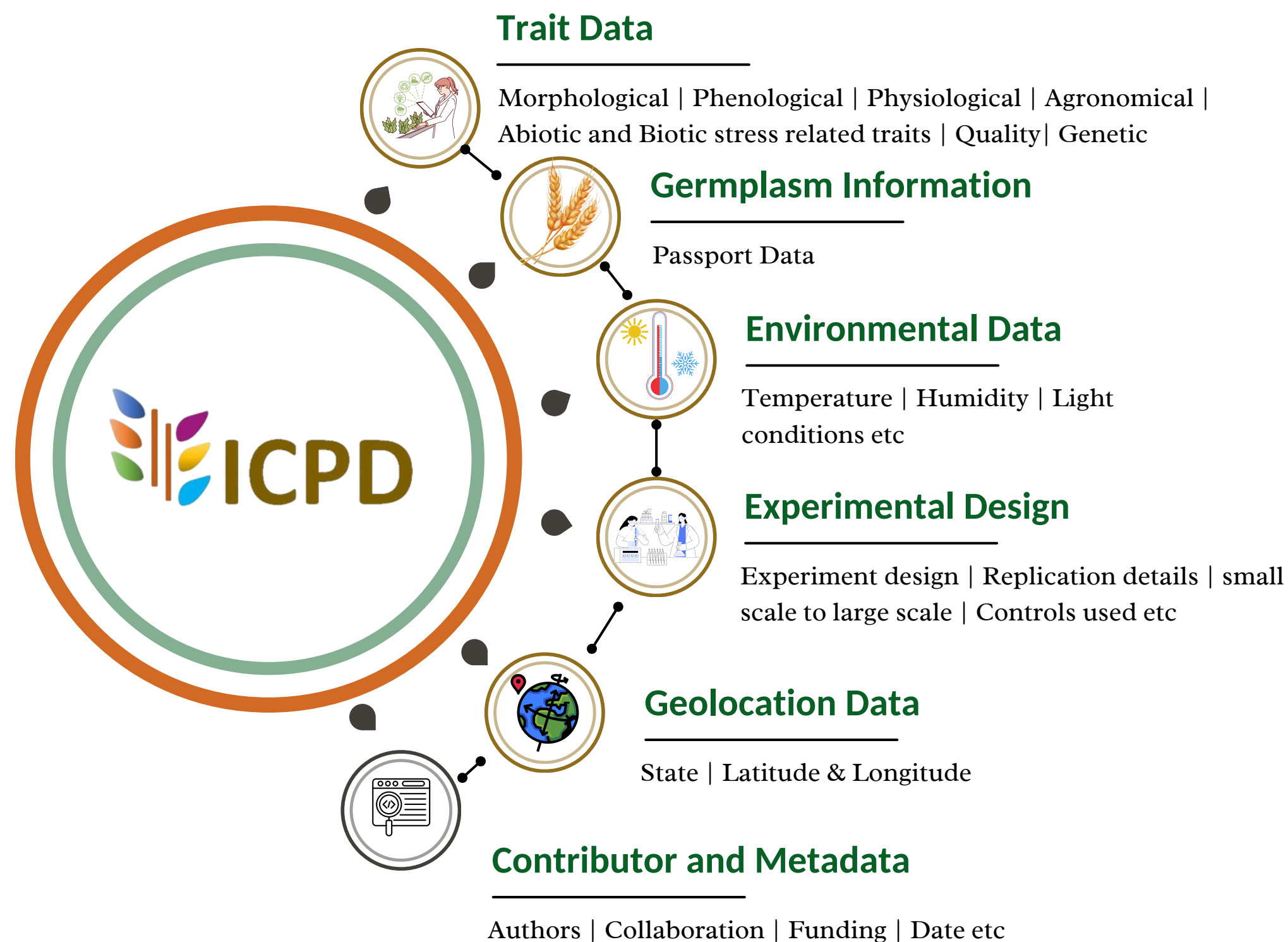
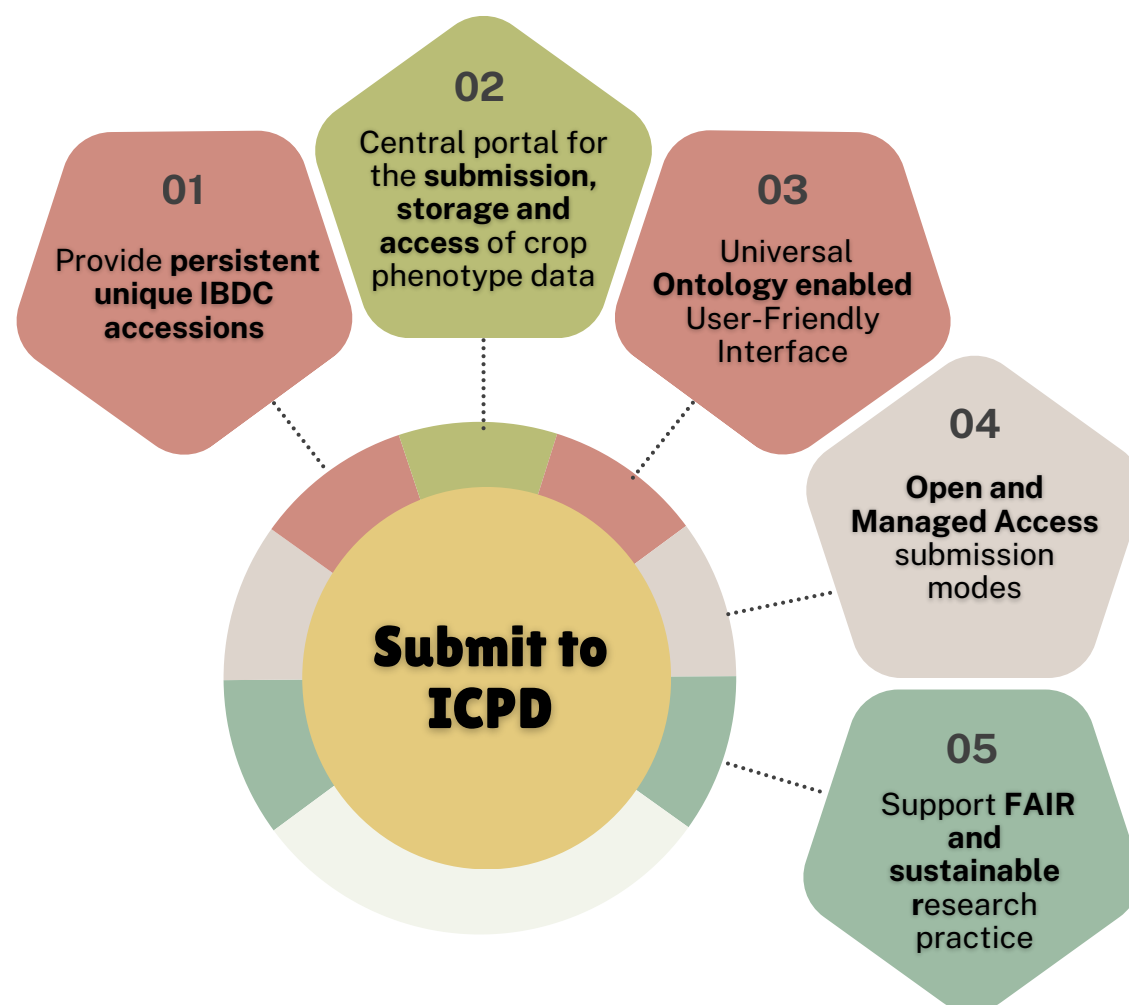
<https://ibdc.dbtindia.gov.in/icpd>

Indian Crop Phenome Database



Vision

- **Biocuration** of crop phenome data: To create a centralized, user-friendly platform for the systemic collection, annotation and integration of phenotype information.
- Aligned with the FAIR (Findable, Accessible, Interoperable, Re-usable) data principles.
- Ensure standardization, accessibility, and interoperability of phenome data through the utilizing structured metadata fields and ontologies to enhance findability, interoperability, and reusability of crop phenomic data, empowering research and development.



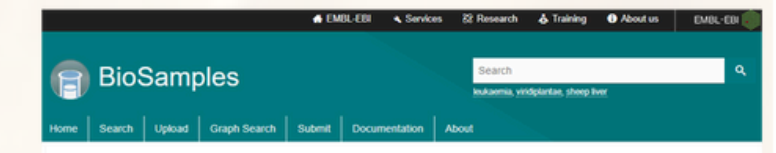
Inspiration



Methods

Enabling reusability of plant phenomic datasets with MIAPPE 1.1

Evangelia A. Papoutsoglou¹, Daniel Faria^{2,3}, Daniel Arend⁴, Elizabeth Arnaud⁵, Ioannis N. Athanasiadis⁶, Ines Chaves^{7,8}, Frederik Coppens^{9,10}, Guillaume Cornut¹¹, Bruno V. Costa^{7,12}, Hanna Cwiek-Kupczyńska¹³, Bert Drosbeke^{9,10}, Richard Finkers¹, Kristina Gruden¹⁴, Astrid Junker⁴, Graham J. King¹⁵, Pawel Krajewski¹⁵, Matthias Lange¹, Marie-Angélique Laporte¹, Célia Michotey¹¹, Markus Oppermann⁴, Richard Ostler¹⁶, Hendrik Poorter^{17,18}, Ricardo Ramirez-Gonzalez¹⁹, Ziva Ramiak¹⁴, Jochen C. Reif⁴, Philippe Rocca-Serra²⁰, Susanna-Assunta Sansone²⁰, Uwe Scholz⁴, François Tardieu²¹, Cristóbal Uauy¹⁹, Björn Usadel^{17,22}, Richard G. F. Visser¹, Stephan Weise⁴, Paul J. Kersey²³, Célia M. Miguel^{7,12}, Anne-Françoise Adam-Blondon¹¹ and Cyril Pommier¹¹

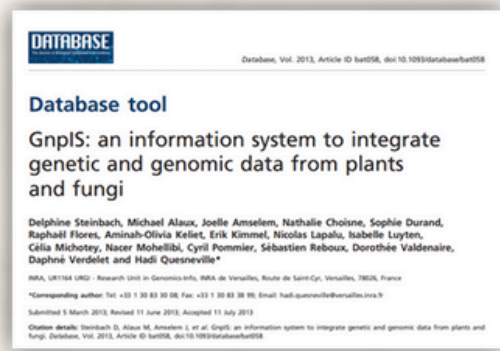
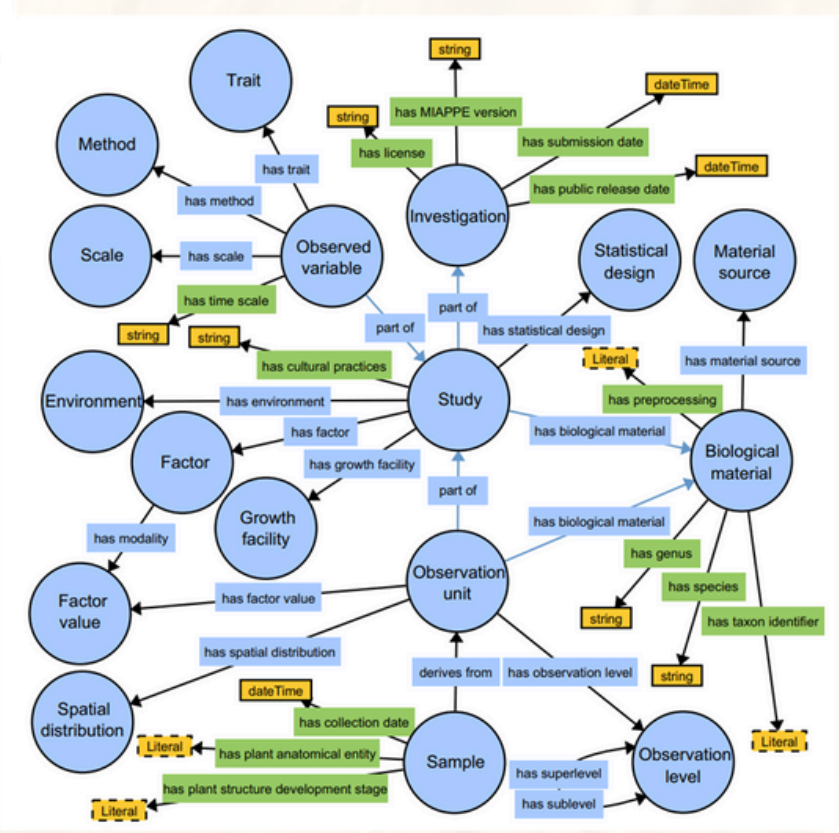


- For plant phenotyping data, **BioSamples** collaborates with the FONDUE project, <https://elixir-europe.org/about-us/commissioned-services/fondue>, implements the Minimum Information About a Plant Phenotyping Experiment (MIAPPE) standards as a sample checklist, and links plant study data in European Nucleotide Archive (ENA) and EVA.

Subset of the Plant Phenotyping Experiment Ontology representing the MIAPPE data model. Generated using WebVOWL (<http://editor.visualdataweb.org/>) and edited manually. Circles indicate classes. Object properties are shown in blue rectangles, and data properties are shown in green rectangles. Yellow rectangles represent literals

MIAPPE

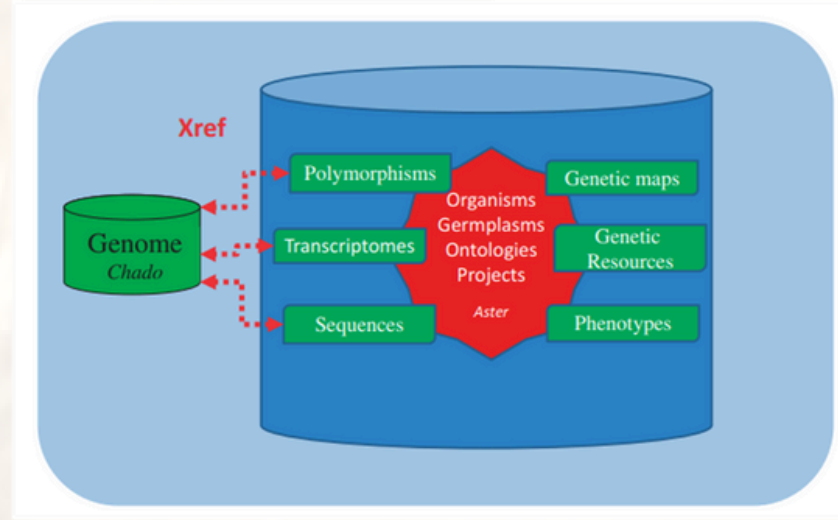
Minimum Information About a Plant Phenotyping Experiment



Genoplante Information System

It bridges genetic and genomic data, allowing researchers access to both genetic information (e.g. genetic maps, quantitative trait loci, markers, single nucleotide polymorphisms, germplasms and genotypes) and genomic data (e.g. genomic sequences, physical maps, genome annotation and expression data) for species of agronomical interest.

- The ‘**phenotype module**’ is dedicated to the storage of phenotype data produced during genotype-environment (climate, soil) interaction studies. Phenotypes can be described with ontologies and linked to data with ‘genetic resources’ and ‘polymorphisms modules’.



GnpIS integrative schema

You are welcome to submit your data into [GnpIS](#). GnpIS handles different types of data in the scope of genetics and genomics for plants including forest trees and fungi: genetic resources, polymorphisms and genotyping data, phenotyping data, association data, genetic maps and QTLs, synteny data.

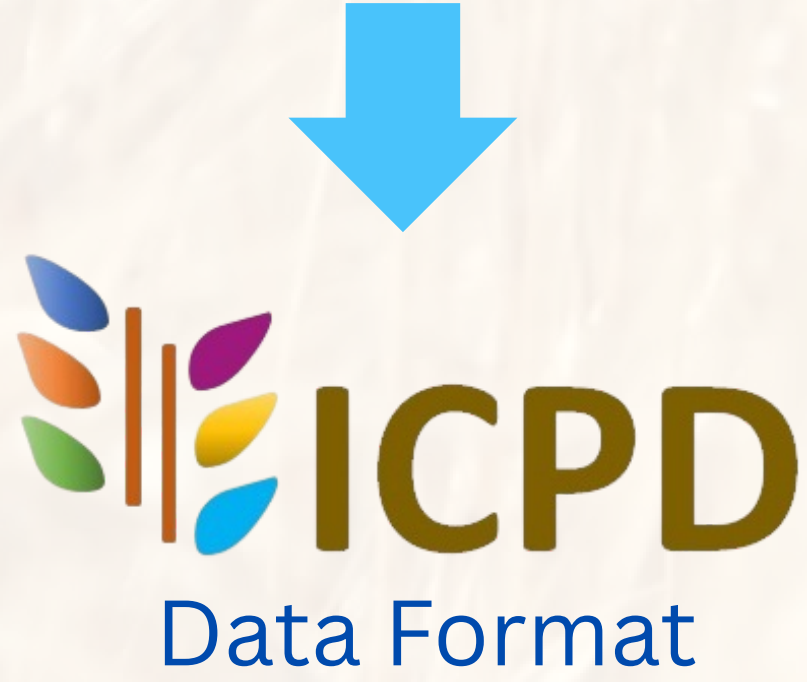
GnpIS gives value to these data by enabling their interoperability and reusability through the use of internationally agreed standards and ontologies: see for instance the [guidelines](#) that were developed in the frame of the WheatIS that are relevant for all species, not only wheat. GnpIS is continuously improving its compliance to the [FAIR](#) principles (Findable, Accessible, Interoperable, Reusable). To enhance the findability and accessibility of the submitted datasets, we collaborate with [data.inra.fr](#) to associate a [DOI](#) (Digital Object Identifier) to your dataset along with its integration in GnpIS.

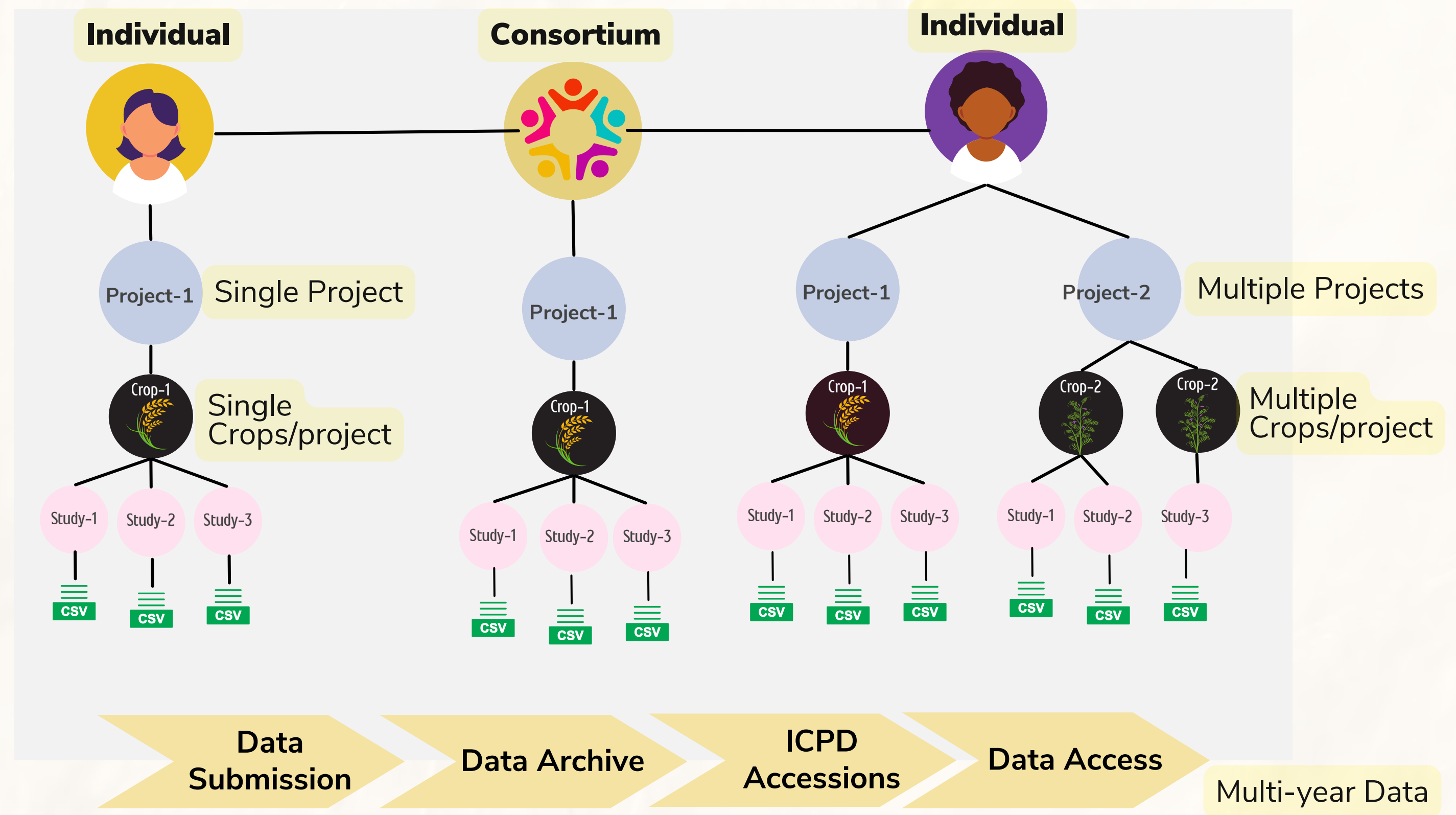
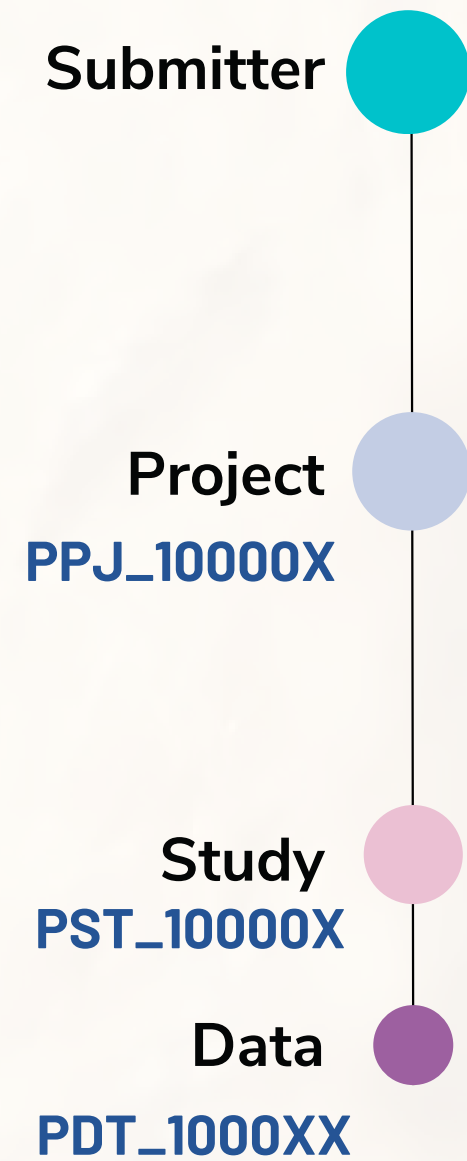
Choose the submission files on the left menu according to the type of data you want to submit to prepare the data. To submit your data into GnpIS, use the data integration [service-offering](#), fill the form and submit it. We will get back to you quickly !

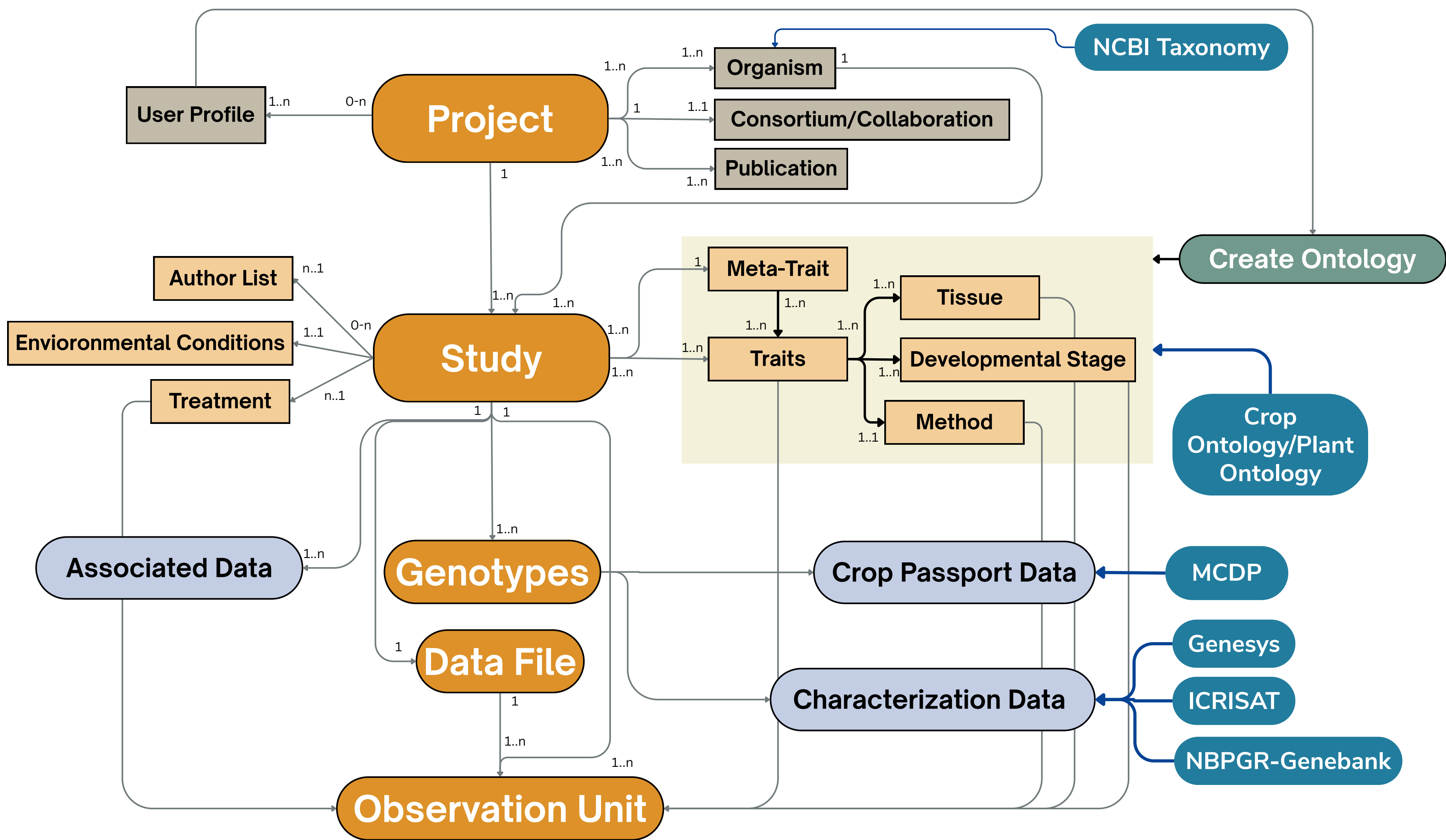
How to submit data :

- **Step 1 :**
 - If the data submission is performed in the frame of an existing collaborative project in partnership with URGI, go directly to step 2.
 - Otherwise, you can submit data through our [service-offering](#) « data insertion and integration in GnpIS », a light procedure allowing to build a minimal data management plan including licence and ownership matters.
- **Step 2 :** Download the template files corresponding to the type of data you want to submit
- **Step 3 :** Fill the template (documentation available)
- **Step 4 :** Submit your files through our [submission form](#)

Different types of files are available according to the type of data :







ICPD VERSION 2 META-DATA MODEL

PHENOPROJECT

Meta Data: Title, Grant Number, Funding Agency, Project Description, Associated Publication, Release date, Collaboration and CROP/s

STUDY

- **Basic Details:** Phenoproject, crop, meta-trait, title of study, description, start and end date, location, experimental design, growth facility, tolerant and susceptible checks
- **Growth & Environment conditions:** Temperature, light, intensity, humidity, etc
- **Treatments:** type (no treatment, biotic, abiotic), treatment agent, description, qualifier, duration and development stage (simultaneous treatment)
- **Traits:** trait name, observation development stage, plant tissue, method details
- **Person:** Authors contributed to the study

DATA FILE TEMPLATE GENERATION

- Select study
- Add accession list of cultivars/genotypes/varieties
- Biological and technical replicates

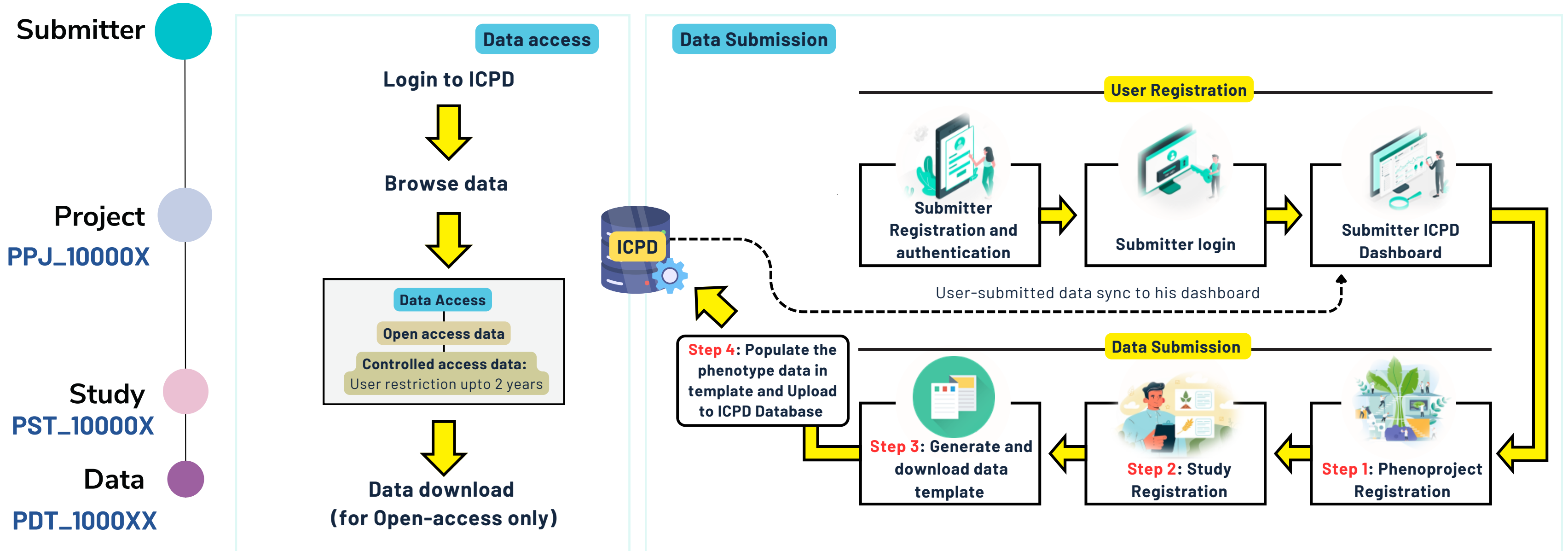
ASSOCIATED DATA SUBMISSION

- Plant Passport data
- Soil and weather data
- Other supporting data



Biological community

Data submission and access flow chart of Indian crop Phenome Database (ICPD)



Data File Format

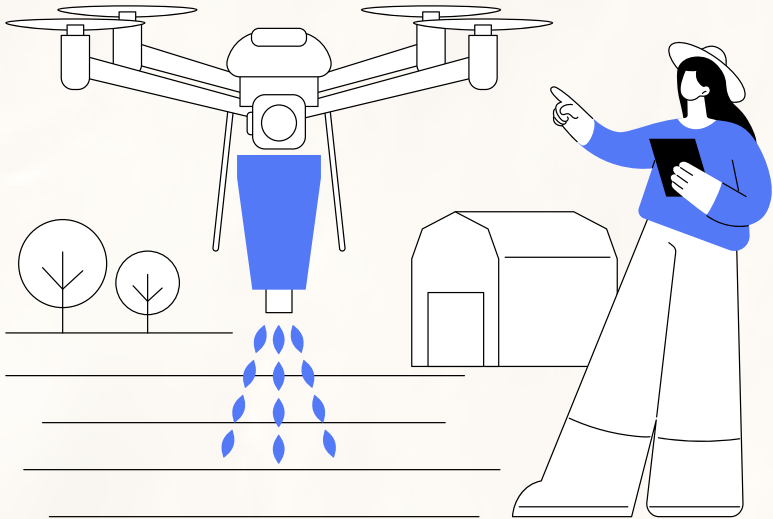
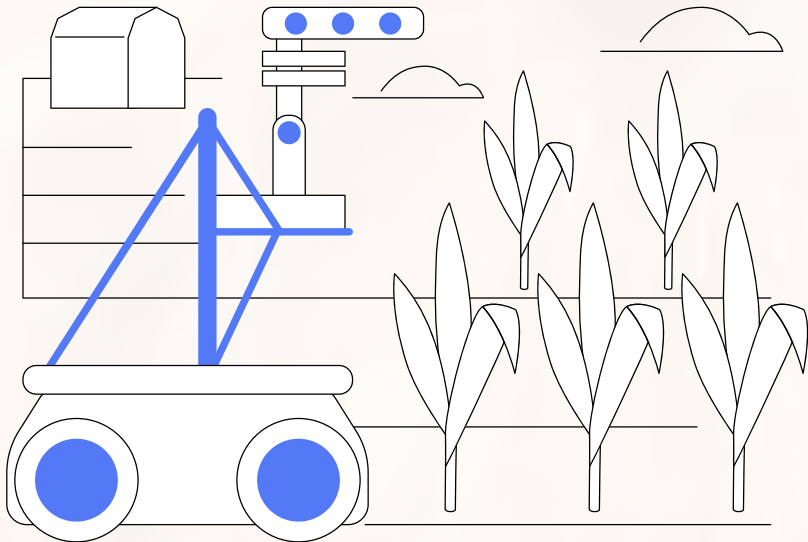
Automatically generated prefilled template with meta-data

Observations

sr_no	study_id	project_id	meta_trait_name	treatment_agent	treatment_qualifier	trait_name	tissue	growthstages	accession	b1_t1	b2_t1	b3_t1	b4_t1	b5_t1	b6_t1	b7_t1	b8_t1	b9_t1	b10_t1
1	PST_100011	PPJ_100010	vegetative vigor			Culm length Rice	vegetative shoot system	40 Day Old Plant	miR408-OE	34.5	33	34	36.5	33.5	35	31	29.5	33	33
2	PST_100011	PPJ_100010	vegetative vigor			Culm length Rice	vegetative shoot system	40 Day Old Plant	WT	27.5	30	27.5	29.5	28.5	29	30	30	31	28
3	PST_100011	PPJ_100010	vegetative vigor			leaf length	leaf	40 Day Old Plant	miR408-OE	25.5	21.5	23.8	20.5	21	20.5	20.7	20	21	22.5
4	PST_100011	PPJ_100010	vegetative vigor			leaf length	leaf	40 Day Old Plant	WT	16.5	22	17	18	18.5	17	16	13	12	17
5	PST_100011	PPJ_100010	vegetative vigor			leaf width	leaf	40 Day Old Plant	miR408-OE	0.45	0.4	0.4	0.4	0.4	0.45	0.4	0.41	0.4	0.45
6	PST_100011	PPJ_100010	vegetative vigor			leaf width	leaf	40 Day Old Plant	WT	0.2	0.2	0.25	0.3	0.3	0.25	0.3	0.25	0.27	0.25
7	PST_100011	PPJ_100010	vegetative vigor			root length	root	40 Day Old Plant	miR408-OE	8.7	8	7.5	9	8	11	9	10	8	6
8	PST_100011	PPJ_100010	vegetative vigor			root length	root	40 Day Old Plant	WT	10	8	5	5	5	6	7	7	9	5
9	PST_100011	PPJ_100010	vegetative vigor			Root density	root	40 Day Old Plant	miR408-OE	19	18	13	20	18	14	19	19	19	15
10	PST_100011	PPJ_100010	vegetative vigor			Root density	root	40 Day Old Plant	WT	13	11	14	13	13	14	13	13	14	14

b = Biological replicates
t = Technical replicates

Template can be used to plan, download and document observations in a structured manner enabling data management and submission to ICPD



ICPD User Dashboard

Browse Ontology

Meta Trait Ontology

Developmental Stage Ontology

Trait Ontology

Plant Tissue Ontology

Method Ontology

List of Available Meta Traits

drought

Search

Search from name or description

MetaTrait ID	Name	Description	Uploaded By
TO_0000467	Cell Membrane Stability	Stability of the cell membrane under the impact of temperature (heat) and water deficit stress.[The trait is often observed in plants as a measure of drought and heat tolerance by determining the amount of solutes/electrolyte leaked from the cell.	
CO_3571000033	Drought Damage Woodyplant	Assessment of drought damages	
CO_34600000269	Drought Induced Senescence Cowpea	Plant senescence due to drought	
CO_32000000060	Drought Injury Rice	The extent to which plant growth is adversely affected by drought. Leaf rolling precedes leaf drying during drought. Response to drought is highly responsive to crop phenology plant growth prior to stress and the timing duration and intensity of drought stress. For many soils it takes at least 2 rainless weeks during the vegetative stage to cause marked differences in response and at least 7 rainless days during the reproductive stage to cause severe drought injury. Repeated ratings are recommended through progress of the drought. Record the stage of plant growth when the stress occurred and the number of stress days.	

Give your feedback

Suggestions

Add your valuable suggestions for improving ICPD services

Technical support

Report here if you encounter any technical error during submission

Create Ontology

Create Meta Trait

Create Developmental Stage

Create Trait

Create Plant Tissue

Create Method

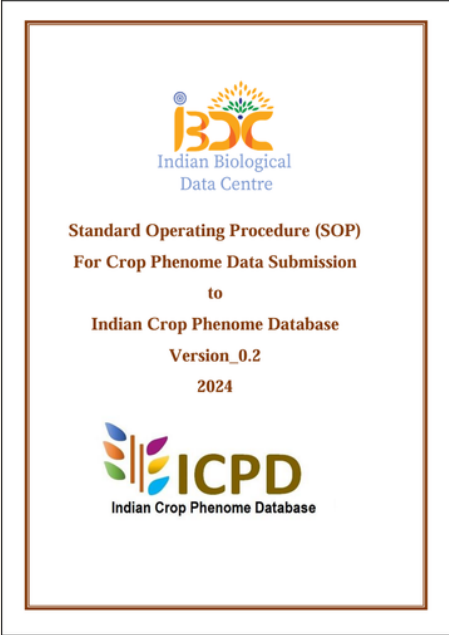
Please ensure that a MetaTrait doesn't already exist before adding your own

Name*

Description*

Submit

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Manage all phenotype projects

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Indian Crop Phenome Database

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Projects Uploaded: 15

Study

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Data Files Uploaded: 12

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Sr.No	Project ID	Project Description	Study ID	Short Description	Upload Date	Modified Date	Uploaded Files
1	PPJ_100072	test1	PST_100054	100 Grain Weight Pearl Millet Greenhouse 2024	Aug. 9, 2024, 11:03 a.m.	Aug. 9, 2024, 11:03 a.m.	View/Download Files
2	PPJ_100072	test1	PST_100053	Drought Induced Senescence Cowpea Greenhouse 2024	Aug. 9, 2024, 10:50 a.m.	Aug. 9, 2024, 10:50 a.m.	View/Download Files
3	PPJ_100051	test_19jan	PST_100052	Abaxial Leaf Vein Pigmentation Sweetpotato Ffghg 2024	Aug. 9, 2024, 10:37 a.m.	Aug. 9, 2024, 10:37 a.m.	No files uploaded
4	PPJ_100070	t	PST_100051	Grain Test Weight Barley Culture Room 2024	Aug. 7, 2024, 11:55 a.m.	Aug. 7, 2024, 11:55 a.m.	View/Download Files
5	PPJ_100070	t	PST_100050	Achene Seed Setting Sunflower Wgg 2024	April 16, 2024, 1:22 p.m.	April 16, 2024, 1:22 p.m.	No files uploaded
6	PPJ_100064	a	PST_100036	10-Dehulled Grain Weight Field 2024	Jan. 25, 2024, 12:11 p.m.	Jan. 25, 2024, 12:11 p.m.	View/Download Files
7	PPJ_100051	test_19jan	PST_100033	10-Dehulled Grain Weight Field 2023	Jan. 19, 2024, 2:10 p.m.	Jan. 19, 2024, 2:10 p.m.	View/Download Files
8	PPJ_100015	q	PST_100031	10-Dehulled Grain Weight Field 2023	Nov. 6, 2023, 12:23 p.m.	Nov. 6, 2023, 12:23 p.m.	View/Download Files
9	PPJ_100014	qq	PST_100030	Abaxial Stomatal Frequency Field 2023	Nov. 3, 2023, 9:33 a.m.	Nov. 3, 2023, 9:33 a.m.	View/Download Files
10	PPJ_100013	test	PST_100025	Abaxial Leaf Vein Pigmentation Sweetpotato Test 2023	Nov. 3, 2023, 9:28 a.m.	Nov. 3, 2023, 9:28 a.m.	View/Download Files
11	PPJ_100008	test	PST_100010	Drought Damage Woodyplant Field 2023	July 27, 2023, 12:13 p.m.	July 27, 2023, 12:13 p.m.	No files uploaded
12	PPJ_100008	test	PST_100009	Grain Filling Rate Rice Fkg 2023	June 6, 2023, 3:06 p.m.	June 6, 2023, 3:06 p.m.	View/Download Files
13	PPJ_100003	Test	PST_100005	10-Dehulled Grain Weight Field 2023	April 19, 2023, 10:25 a.m.	April 19, 2023, 10:25 a.m.	View/Download Files
14	PPJ_100003	Test	PST_100004	Grain Test Weight Barley Field 2023	April 18, 2023, 4:36 p.m.	April 18, 2023, 4:36 p.m.	View/Download Files

See and download the metadata of project and study

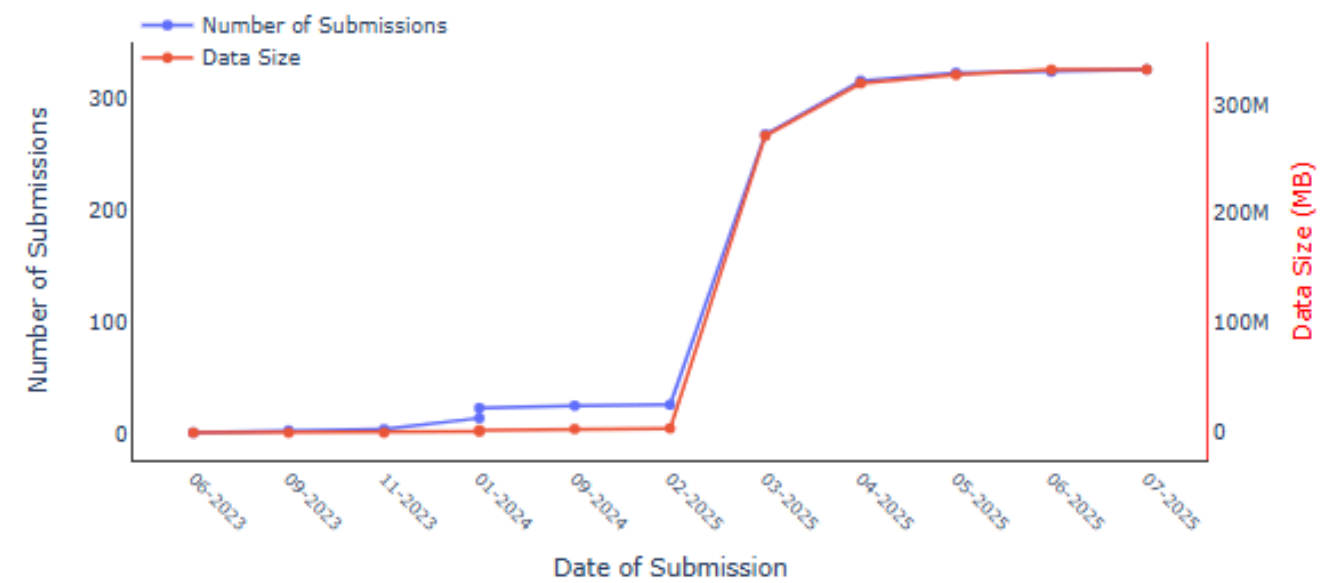
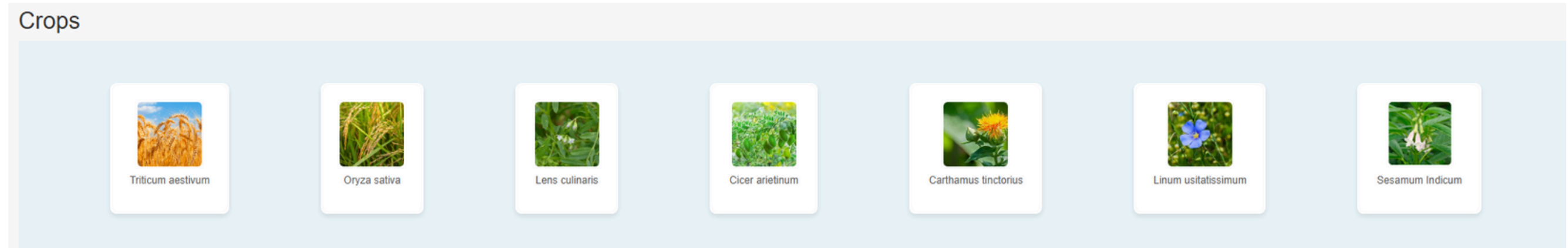
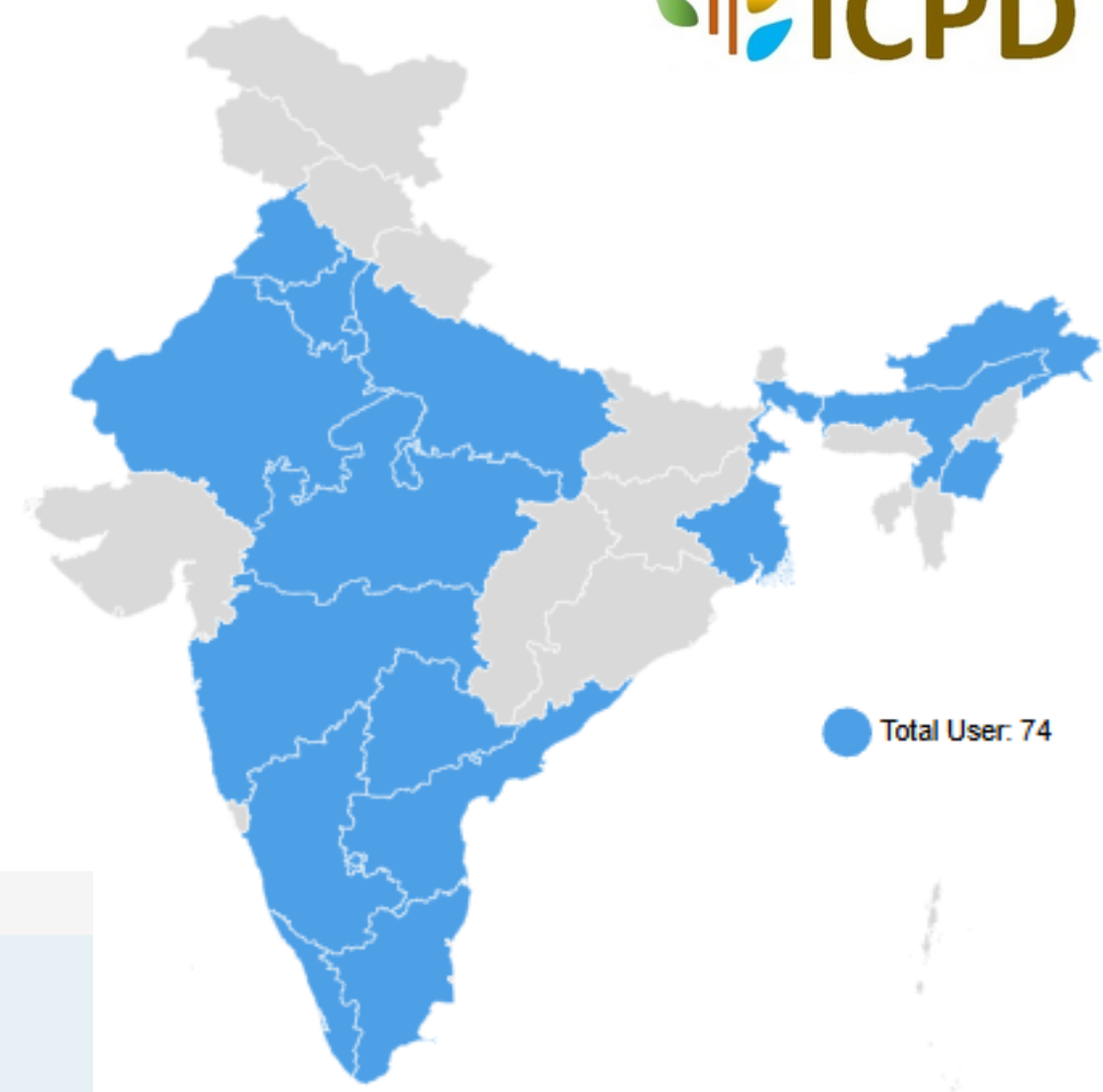
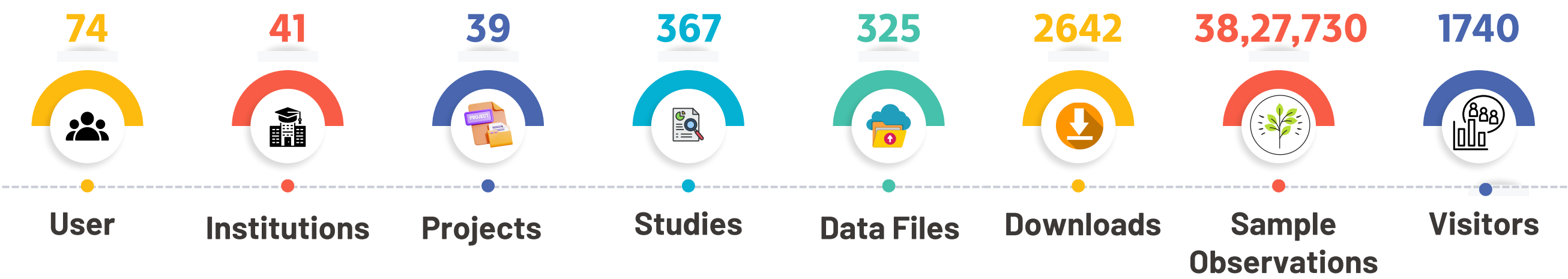
Details

Timestamp

CURRENT DATA STATS

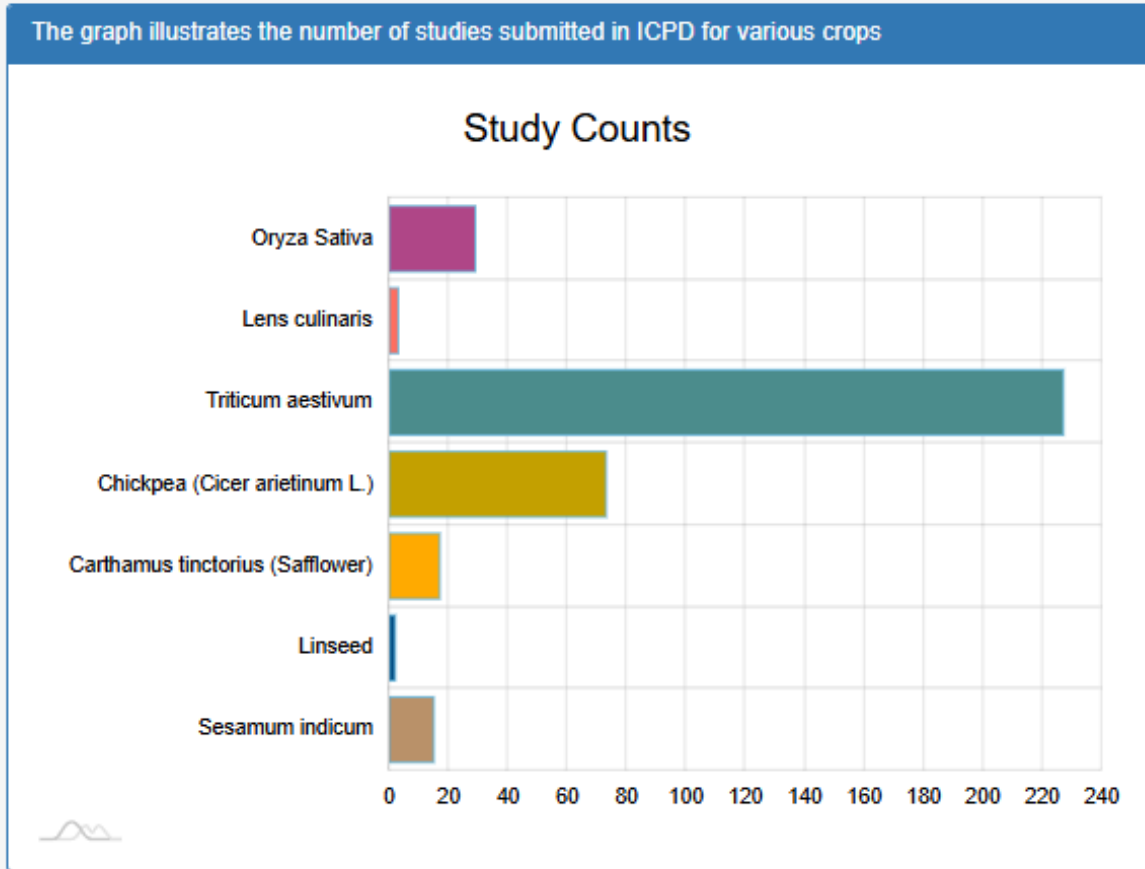


Submitted Data Stats

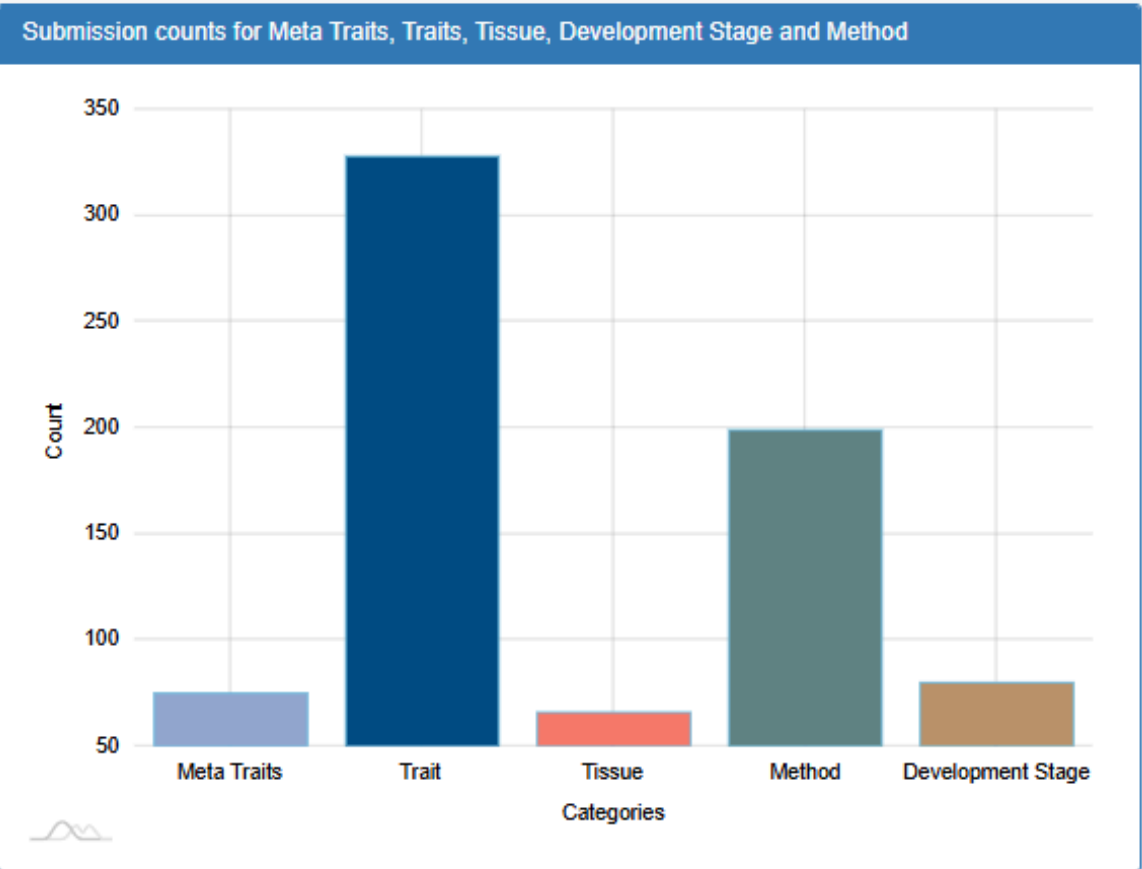


CURRENT DATA STATS

Study Distribution by Crops



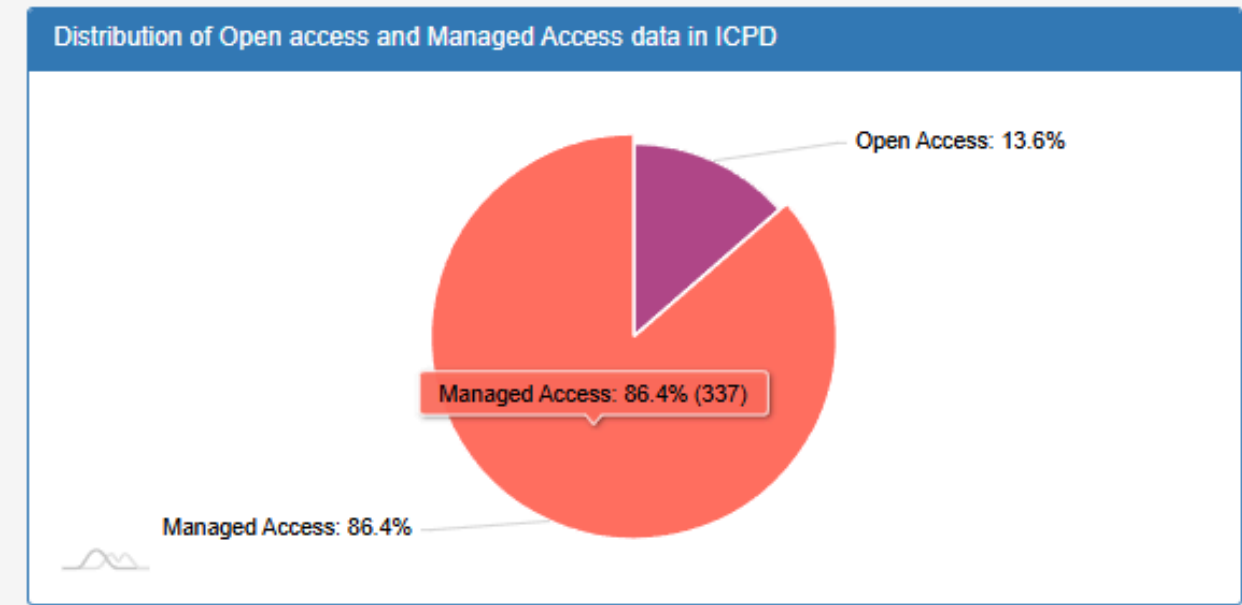
Trait Details



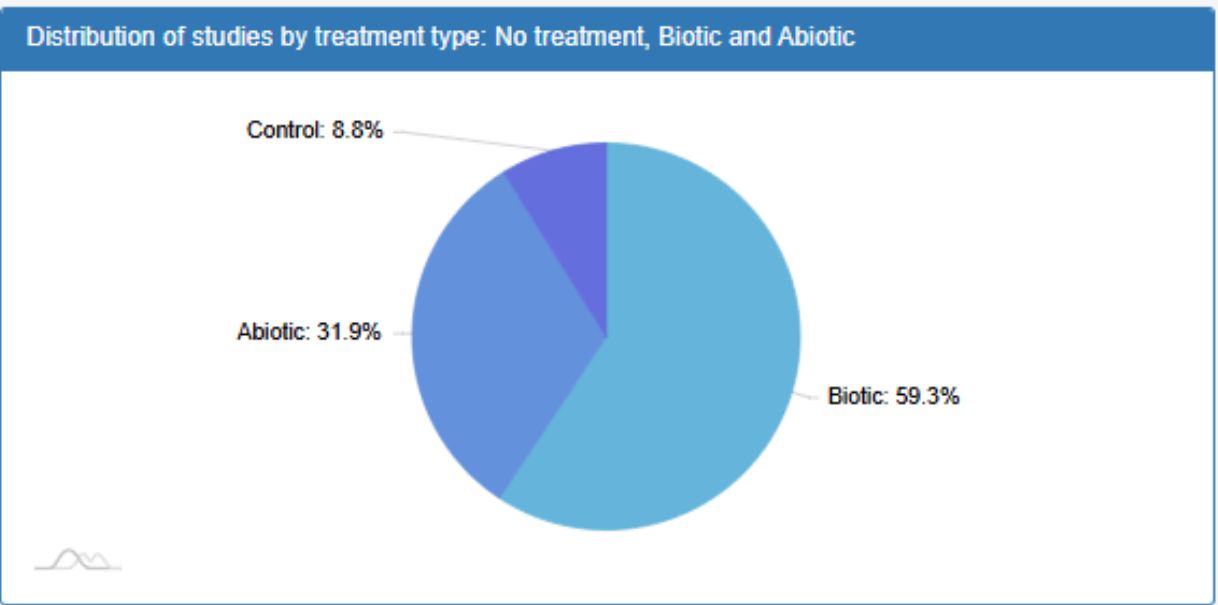
37 Meta Traits



Data Access

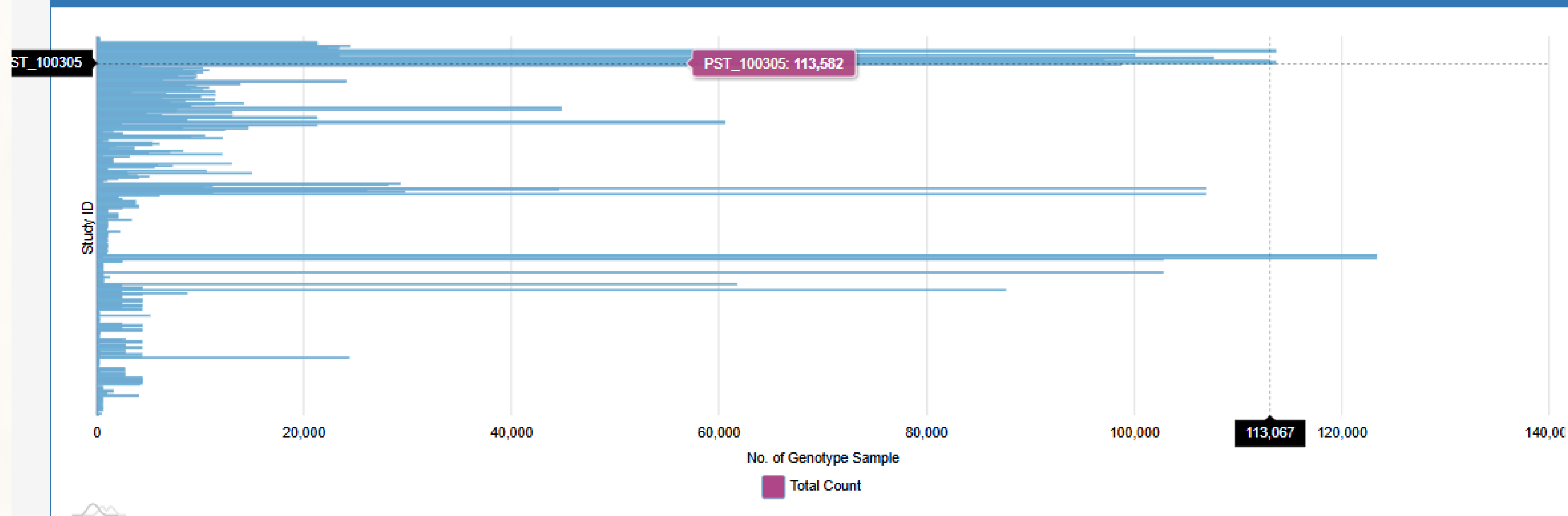


Treatment Type



Genotype Samples

Distribution of Genotype Samples per study



ICPD is the data hub for major mission-mode programs on "Characterization of Genetic Resources," supported by the DBT

Major Mission Projects

Triticum aestivum



Germplasm Characterization and Trait Discovery in Wheat using Genomics Approaches and its Integration for Improving Climate Resilience, Productivity and Nutritional quality.

Oryza sativa



Mainstreaming rice landraces diversity in varietal development through genome-wide association studies: A model for large-scale utilization of gene bank collections of rice.

Linum usitatissimum



Leveraging genetic resources for accelerated genetic improvement of Linseed using comprehensive genomics and phenotyping approaches.

Cicer arietinum



Characterization of Chickpea Germplasm Resource to Accelerate Genomics-assisted Crop Improvement.

Carthamus tinctorius



Exploiting Genetic Diversity for Improvement of Safflower through Genomics Assisted Discovery of QTLs/Genes Associated with Agronomic Traits.

Sesamum indicum



Mainstreaming sesame germplasm for productivity enhancement and sustainability through genomics assisted core development and trait discovery.

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Chickpea Atlas for National Agri-innovation

Characterization of Chickpea Germplasm Resource to Accelerate Genomics-assisted Crop Improvement

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ABOUT

A multi-institutional National Mission Program on "Characterization of Chickpea Germplasm Resource to Accelerate Genomics-assisted Crop Improvement" funded by Department of Biotechnology (DBT), Ministry of Science and Technology, Govt. of India has been launched in the year 2020 involving 15 participating institutions from Indian Council of Agricultural Research (ICAR), state agricultural universities, and international organization in which Biotechnology Research and Innovation Council-National Institute of Plant Genome Research (BRIC-NIPGR) serves as a Coordinating partner. The Mission Project envisages comprehensive phenotypic and genotypic characterization of a Pan Genebank Core of 5084 germplasm accessions (ICRISAT and NBPGR) and 6004 trait-specific genotypes, including germplasm accessions, pre-breeding lines, genetic stocks, wild Cicer accessions and released varieties, to identify novel superior genes/alleles governing traits of agronomic importance and promising trait-specific accessions (donors) for accelerating genomics-assisted breeding and crop improvement in chickpea. To constitute a morphological core, the multi-location DUS agro-morphological characterization of 5084 chickpea germplasm accessions using 22 Descriptor traits has been performed to identify accessions with superior yield component and plant architectural traits. The genotypic characterization of 11088 accessions using whole genome resequencing-based SNP genotyping has been accomplished to constitute a molecular core and subsequently an iterative core of chickpea. Comprehensive phenotypic characterization of 5084 Pan Genebank Core germplasm accessions identified promising accessions resistant/tolerant to the six biotic (Fusarium wilt, Ascochyta blight, Botrytis grey mould, collar rot, dry root rot and nematode resistance) and four abiotic (drought, heat, cold and salinity) stress tolerance traits as well as accessions of erect plant architecture types with superior agronomic performance and amenable to





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Phenotype Genotype Associated Data

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Accession	Institutes
ICC 2	ICRISAT
ICC 12	ICRISAT
ICC 15	ICRISAT
ICC 18	ICRISAT
ICC 26	ICRISAT
ICC 42	ICRISAT
ICC 43	ICRISAT
ICC 45	ICRISAT
ICC 64	ICRISAT
ICC 65	ICRISAT
ICC 66	ICRISAT
ICC 67	ICRISAT
ICC 68	ICRISAT
ICC 75	ICRISAT
ICC 80	ICRISAT
ICC 86	ICRISAT

Phenotype Genotype Associated Data

Sr.No.	Data type	Repository Name	Study accession	Study title	Run count	Centre Name	Date of submission
1	Whole Genome Sequencing	INDA-CA	INCARP000305	Genomic Characterization of Chickpea Pan-genebank Genetic Resources	10264	National Institute of Plant Genome Research, NIPGR, New Delhi	28-04-2025

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Phenotype Genotype Associated Data

Total submitted Files: 56

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Search:

Study Accession	Study Title	Release date	Study Submitted Date	Crop	Traits	Institute	Data Size	Data file
PST_100081	Screening of chickpea germplasm for resistance to dry root rot disease	31-03-2027	08-03-2025	Cicer arietinum L.	Dry root rot resistance Chickpea	National Institute of Plant Genome Research	708.63 KB	Control Access
PST_100122	DUS agromorphological trait characterization of chickpea pan-genebank core accessions	31-03-2027	12-03-2025	Chickpea (Cicer arietinum L.)	DUS_AGROMORPHOLOGY_TRAITS_CHICKPEA	National Institute of Plant Genome Research	676.4 KB	Control Access
PST_100141	DUS agromorphological trait characterization of chickpea pan-genebank core accessions	31-03-2027	13-03-2025	Chickpea (Cicer arietinum L.)	DUS_AGROMORPHOLOGY_TRAITS_CHICKPEA	National Institute of Plant Genome Research	14.08 MB	Control Access
PST_100173	Screening of chickpea accessions for resistance to Fusarium wilt (causal agent : Fusarium oxysporum f. sp. ciceri)	31-03-2027	21-03-2025	Chickpea (Cicer arietinum L.)	Fusarium wilt resistance Chickpea	National Institute of Plant Genome Research	562.91 KB	Control Access
PST_100173	Screening of chickpea accessions for resistance to Fusarium wilt (causal agent : Fusarium oxysporum f. sp. ciceri)	31-03-2027	21-03-2025	Chickpea (Cicer arietinum L.)	Fusarium wilt resistance Chickpea	National Institute of Plant Genome Research	562.91 KB	Control Access
PST_100329	Screening of chickpea germplasm for drought stress tolerance	31-03-2027	02-04-2025	Chickpea (Cicer arietinum L.)	drought	National Institute of Plant Genome Research	542.35 KB	Control Access

Accession Details: IC75500

Note:
Data Source: Characterization data for Chickpea has been obtained from the ICRISAT Genebank. [ICRISAT Chickpea Characterization Data.](#)
[← Back to Accessions](#)

Passport Information

Accession Identifier: IC75500
Institutes: NBPGR
ICRISAT_NBPGR Identifier: ICAR-NBPGR
Crop: Chickpea
DOI: NA
Mission_code: NA
Collectors_number: NA
Alternate_accession_identifier: E-235
Local Name: NA
Genus: Cicer
Species: arietinum
Sp_author: NA
Subtaxa: NA
Subauthor: NA
Cultivar Name: NA
Biological Status: NA
Cultivar Types: NA
Collecting Source: NA
Donor Cooperator_code: NA
Donor Country: NA
Acquisition Date: NA
Collection Date: NA
Country Source: India
Province: Unknown
Collection Site: NA
Latitude: NA
Longitude: NA
Elevation: NA
EC_No: NA
FAO in Trust: NA
Core Collection: NA
Mini Core Collection: NA
Collectors Number Repeated: NA

Characterization Information

Growth Habit:
Plant Height (cm):
Plant Width (cm):
Flower Color:
Days to Flowering:
Days to Maturity:
Pods per Plant:
Seeds per Pod:
Seed Shape:
Seed Surface:
100 Seed Weight (g):
Seed Yield (kg/ha):
Protein (%):
Project:

Ontology enabled data submission



Browse Ontology

Meta Trait Ontology

Developmental Stage Ontology

Trait Ontology

Plant Tissue Ontology

Method Ontology

List of Available Meta Traits

drought

Search

Search from name or description

MetaTrait ID	Name	Description	Uploaded By
TO_0000467	Cell Membrane Stability	Stability of the cell membrane under the impact of temperature (heat) and water deficit stress.[The trait is often observed in plants as a measure of drought and heat tolerance by determining the amount of solutes/electrolyte leaked from the cell.	
CO_357:1000033	Drought Damage Woodyplant	Assessment of drought damages	
CO_340:0000269	Drought Induced Senescence Cowpea	Plant senescence due to drought	
CO_320:0000060	Drought Injury Rice	The extent to which plant growth is adversely affected by drought. Leaf rolling precedes leaf drying during drought. Response to drought is highly responsive to crop phenology plant growth prior to stress and the timing duration and intensity of drought stress. For many soils it takes at least 2 rainless weeks during the vegetative stage to cause marked differences in response and at least 7 rainless days during the reproductive stage to cause severe drought injury. Repeated ratings are recommended through progress of the drought. Record the stage of plant growth when the stress occurred and the number of stress days.	

Create Ontology

Create Meta Trait

Create Developmental Stage

Create Trait

Create Plant Tissue

Create Method

Please ensure that a MetaTrait doesn't already exist before adding your own

Name*

Description*

Submit

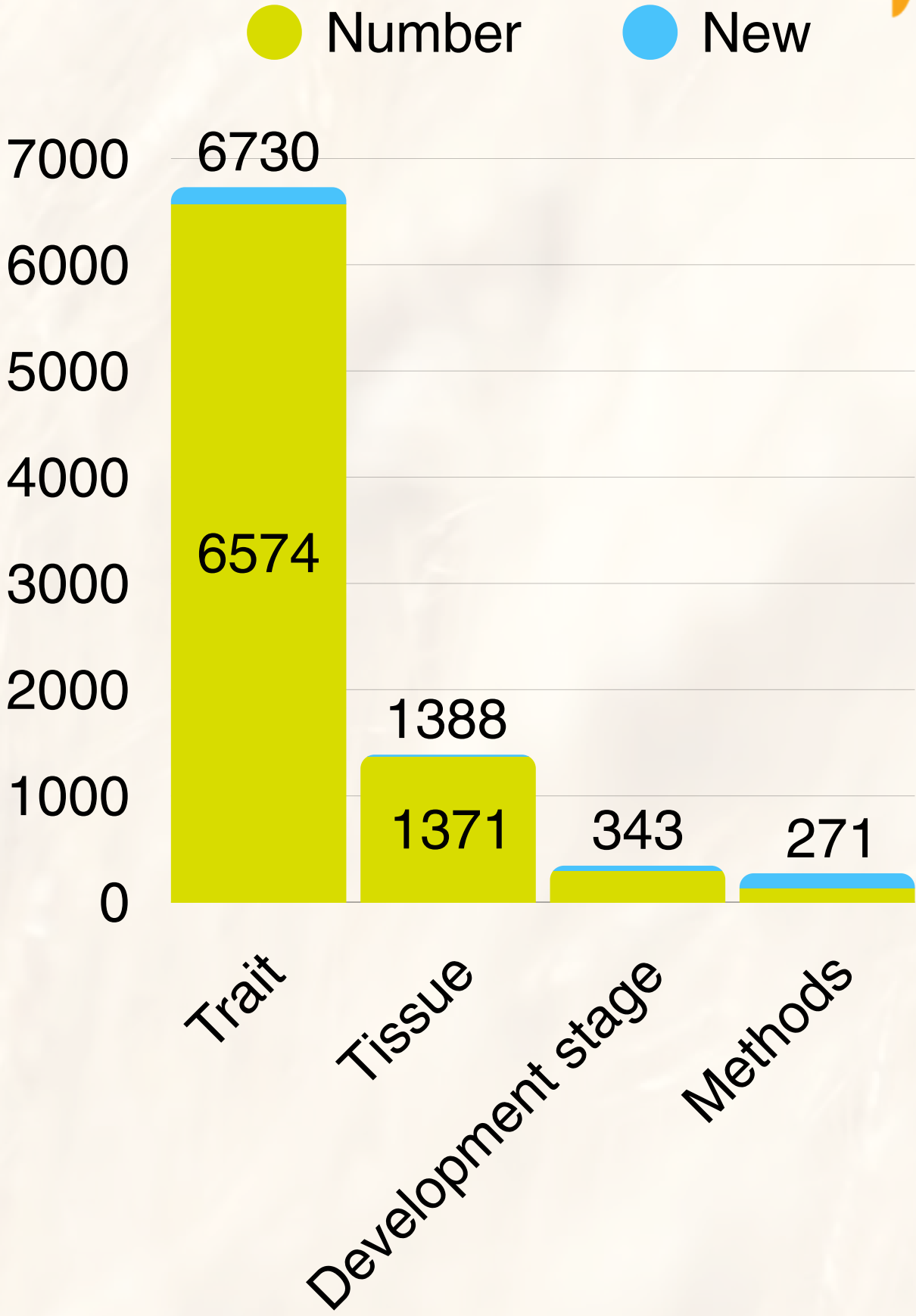
MCMT106562 (Meta Trait)

MCT106562 (Trait)

MCGS100297 (Stage)

MCPT101371 (Plant Tissue)

MCM100130 (Method)



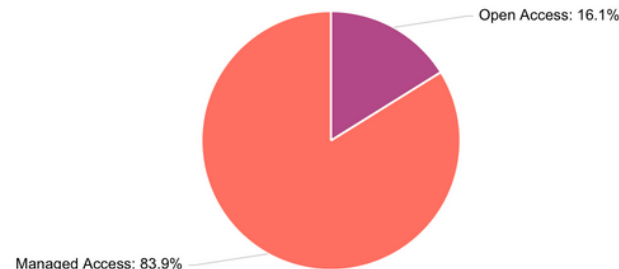
Submitter can contribute to ontology creation



Browse Data: Open and managed access

Data Access

Distribution of Open access and Managed Access data in ICPD



Total submitted Files: 196

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Search:

Phenoproject Accession	Phenoproject Title	Submitted Date	Release date	Study Accession	Study Title	Crop	Traits	Institute	Data Size	Data file
PPJ_100069	"Mainstreaming rice landraces diversity in varietal development through genome wide association studies: A model for large-scale utilization of gene bank collections of rice"	30-01-2024	03-11-2023	PST_100048	Screening of rice varieties for resistance to brownplanthopper (Causal agent: Nilparvata lugens)	Oryza Sativa	brown planthopper resistance	Acharya N. G. Ranga Agricultural University, RARS	82.06 KB	View/Download Files
PPJ_100069	"Mainstreaming rice landraces diversity in varietal development through genome wide association studies: A model for large-scale utilization of gene bank collections of rice"	30-01-2024	03-11-2023	PST_100049	Screening of rice varieties for resistance to leaf blast (Causal agent: Magnaporthe grisea)	Oryza Sativa	leaf blast disease resistance	Acharya N. G. Ranga Agricultural University, RARS	169.87 KB	View/Download Files
PPJ_100069	"Mainstreaming rice landraces diversity in varietal development through genome wide association studies: A model for large-scale utilization of gene bank collections of rice"	30-01-2024	03-11-2023	PST_100059	Screening of rice varieties for resistance to leaf blast (Causal agent: Magnaporthe grisea)	Oryza Sativa	panicle blast disease resistance	Acharya N. G. Ranga Agricultural University, RARS	169.21 KB	View/Download Files
PPJ_100069	"Mainstreaming rice landraces diversity in varietal development through genome wide association studies: A model for large-scale utilization of gene bank collections of rice"	30-01-2024	03-11-2023	PST_100064	Screening of rice varieties for resistance to brown planthopper (Causal agent: Nilparvata lugens)	Oryza Sativa	brown planthopper resistance	Acharya N. G. Ranga Agricultural University, RARS	82.89 KB	View/Download Files
PPJ_100069	"Mainstreaming rice landraces diversity in varietal development through genome wide association studies: A model for large-scale utilization of gene bank collections of rice"	30-01-2024	03-11-2023	PST_100200	Screening of rice varieties for resistance to bacterial blight	Oryza Sativa	rice bacterial blight disease resistance	Acharya N. G. Ranga Agricultural University, RARS	111.48 KB	View/Download Files
PPJ_100075	Characterization of a genetically diverse reference set of lentil	23-09-2024	21-09-2026	PST_100057	Estimation of seed mineral content in a genetically diverse reference set of lentil	Lens culinaris	mineral and ion content trait	National Institute of Plant Genome Research (NIPGR), Delhi	482.16 KB	Control Access
PPJ_100075	Characterization of a genetically diverse reference set of lentil	23-09-2024	21-09-2026	PST_100058	Estimation of seed mineral content in a genetically diverse reference set of lentil (2023-24)	Lens culinaris	mineral and ion content trait	National Institute of Plant Genome Research (NIPGR), Delhi	486.85 KB	Control Access
PPJ_100086	Germplasm Characterization and Trait Discovery in Wheat using Genomics Approaches and its Integration for Improving Climate Resilience, Productivity and Nutritional quality	24-02-2025	23-02-2027	PST_100070	Characterization of wheat germplasm lines for stripe rust resistance at SKUAST, Kashmir (2020-21)	Triticum aestivum	wheat stripe rust disease resistance	ICAR-National Nureau of Plant Genetic Resources	561.71 KB	Control Access
PPJ_100086	Germplasm Characterization and Trait Discovery in Wheat using Genomics Approaches and its Integration for Improving Climate Resilience, Productivity and Nutritional quality	24-02-2025	23-02-2027	PST_100071	Characterization of wheat germplasm lines for stripe rust resistance at SKUAST, Kashmir (2020-21)	Triticum aestivum	wheat stripe rust disease resistance	ICAR-National Nureau of Plant Genetic Resources	590.67 KB	Control Access
PPJ_100086	Germplasm Characterization and Trait Discovery in Wheat using Genomics Approaches and its Integration for Improving Climate Resilience, Productivity and Nutritional quality	24-02-2025	23-02-2027	PST_100072	Characterization of wheat germplasm lines for stripe rust resistance at SKUAST, Kashmir (2021-22)	Triticum aestivum	wheat stripe rust disease resistance	ICAR-National Nureau of Plant Genetic Resources	590.67 KB	Control Access

Showing 21 to 30 of 196 entries

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Data Sharing

Study Details

Project

Project uploaded by:	RARS MTU ,principal_investigator , Acharya N. G. Ranga Agricultural University, RARS
Created:	30-01-2024
Project ID:	PPJ_100069
Project Title:	"Mainstreaming rice landraces diversity in varietal development through genome wide association studies: A model for large-scale utilization of gene bank collections of rice"
Project Description:	Phenotyping of rice landraces for biotic stresses BPH, BLB and BLAST
Grant Number:	239
Funding Agency:	DBT
Associated Publications (Title/DOI/PMID):	NA
Release Date:	Nov. 3, 2023
Project Type:	Consortium

Study

Study uploaded by:	RARS
Created:	20-02-2024
Study ID:	PST_100048
Crop ID:	Oryza Sativa
Meta-Trait:	brown planthopper resistance
Study Title:	Screening of rice varieties for resistance to brownplanthopper (Causal agent: Nilparvata lugens)
Description:	
Start Date :	June 30, 2022
End Date:	Nov. 14, 2022
Data Type:	Phenotypic
Location :	RARS, Maruteru
Description of experimental design :	single observational plot
Growth Facility :	field
Tolerent Check :	PTB 33
Susceptible Check:	TN1

Growth and Environmental Conditions

Temperature Lower Limit (°C):	26.000
Temperature Upper Limit (°C):	31.000
Light Intensity Lower Limit (lux):	0.000
Light Intensity Upper Limit (lux):	0.000
Relative Humidity Lower Limit (%):	67.000
Relative Humidity Upper Limit (%):	89.000
Other Conditions:	

Treatments

Sr.No	Treatment Type	Treatment Developmental Stage	Treatment Name	Treatment Value	Treatment Duration	Treatment Description
1	biotic	plant tissue development stage PO:0025423	Nilaparvatha Lugens	Biotype 4	4 months	Natural infestation of brown planthopper under field conditions

Traits

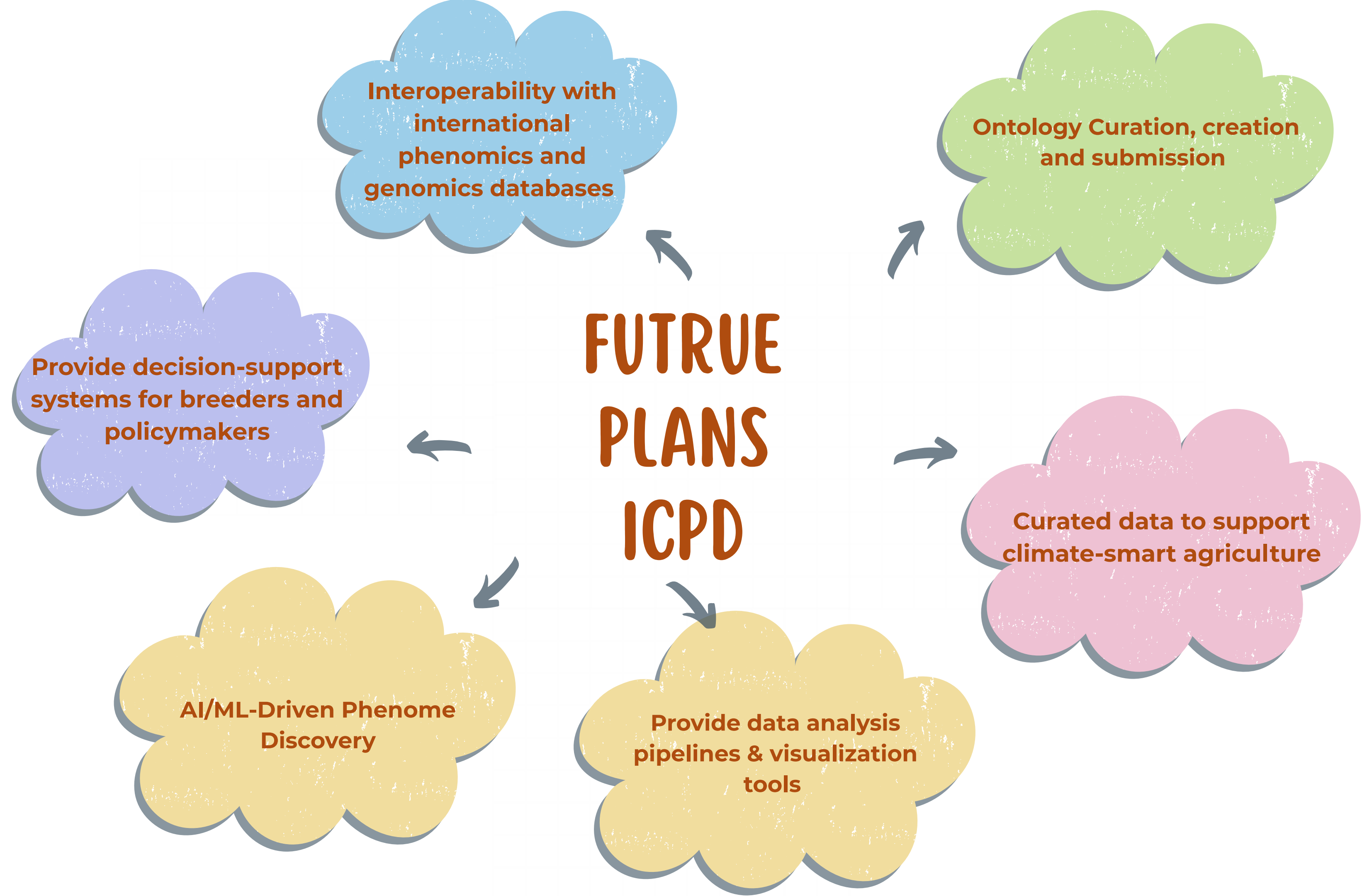
Sr.No	Trait Name	Observation Developmental Stage	Observed Plant Tissue	Method Name
1	brown planthopper resistance TO_0000424	whole plant development stage PO:0007033	whole plant PO:0000003	brown planthopper injury assessment field brown planthopper injury scale field Ordinal

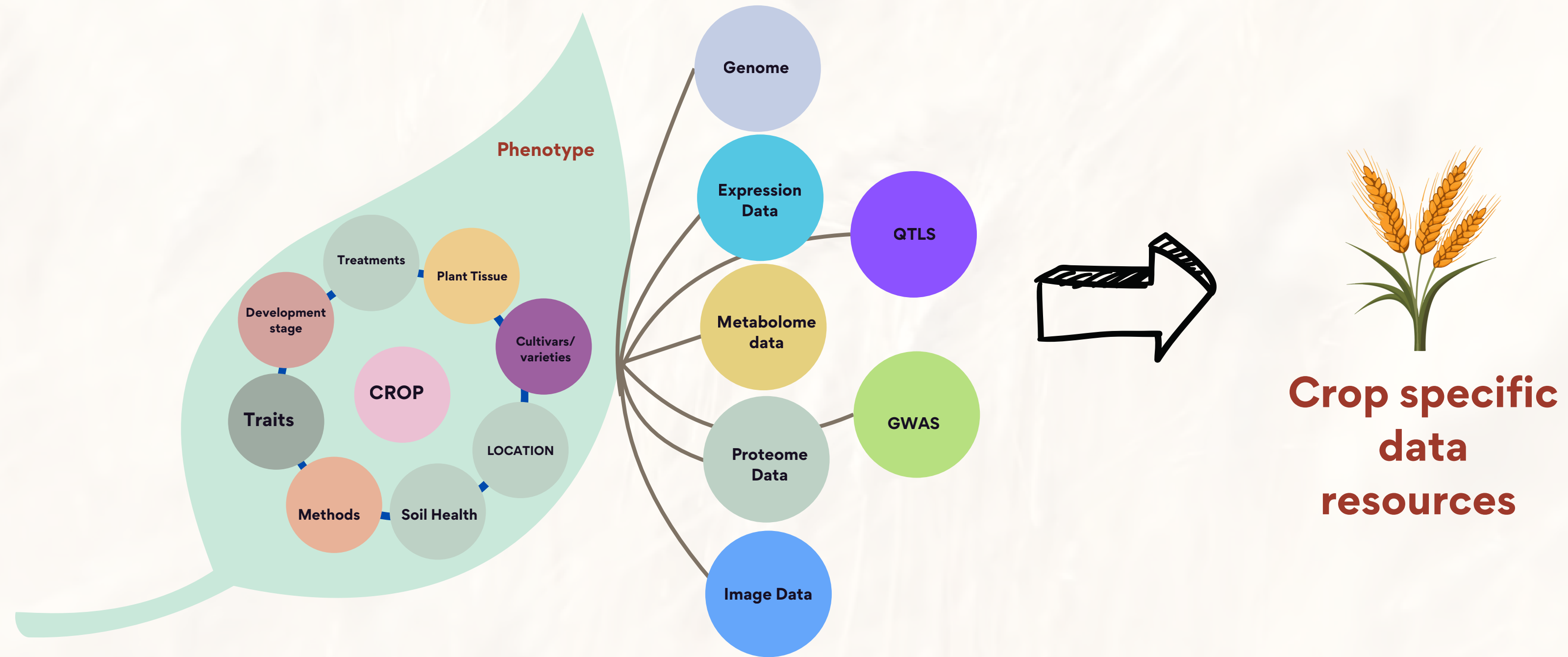
Author Details

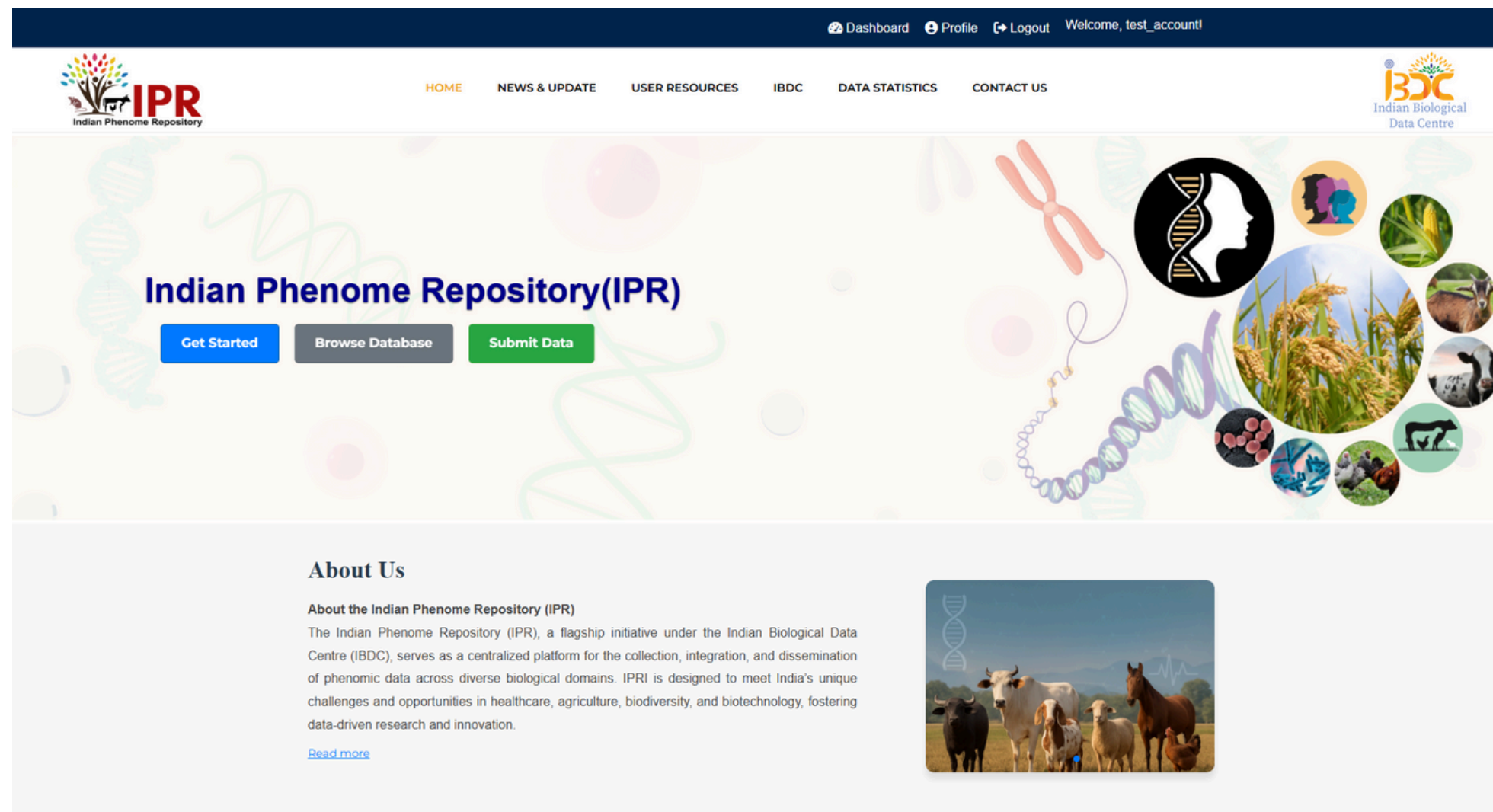
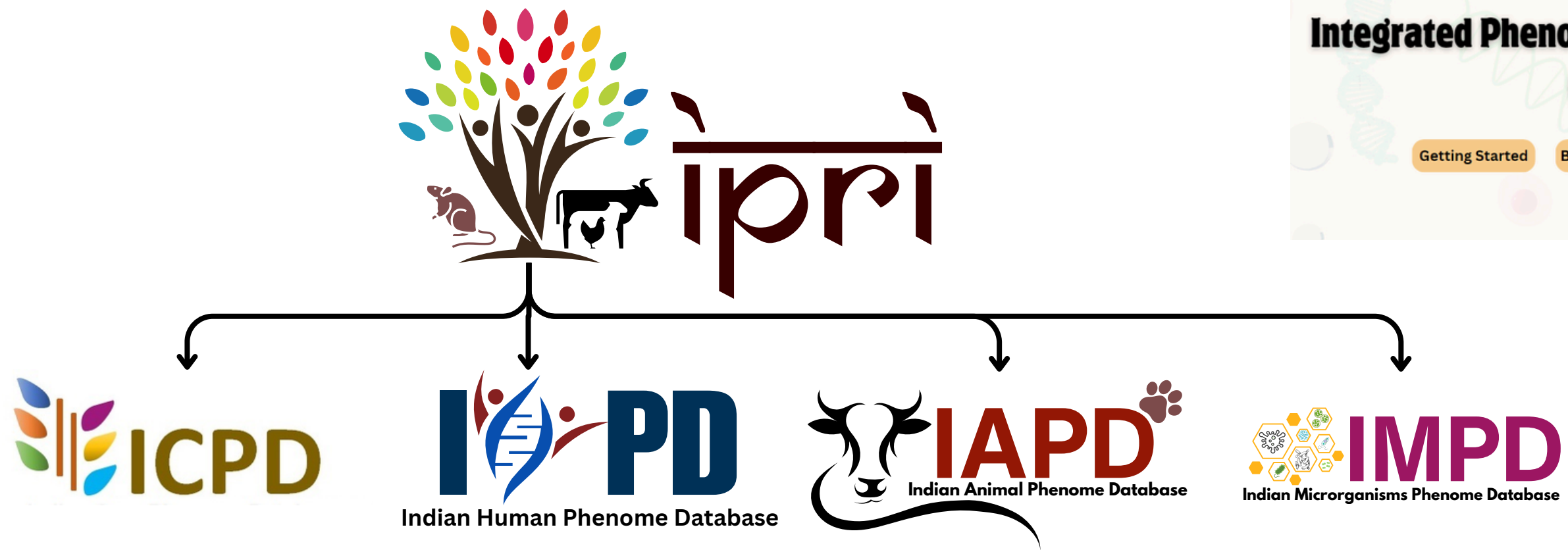
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6	Bhuvaneswari	V	v.bhuvaneswari@angrau.ac.in	Acharya N. G. Ranga Agricultural University, RARS	co_principal_investigator
7	Ravi Kumar	BNVSR	bnvsr.ravikumar@angrau.ac.in	Acharya N. G. Ranga Agricultural University, RARS	principal_investigator

Uploaded Files

Sr.No.	Data File Accession	No. of Genotype Samples	Size of File	Download
1	PDT_100048	500	82.1 KB	Download

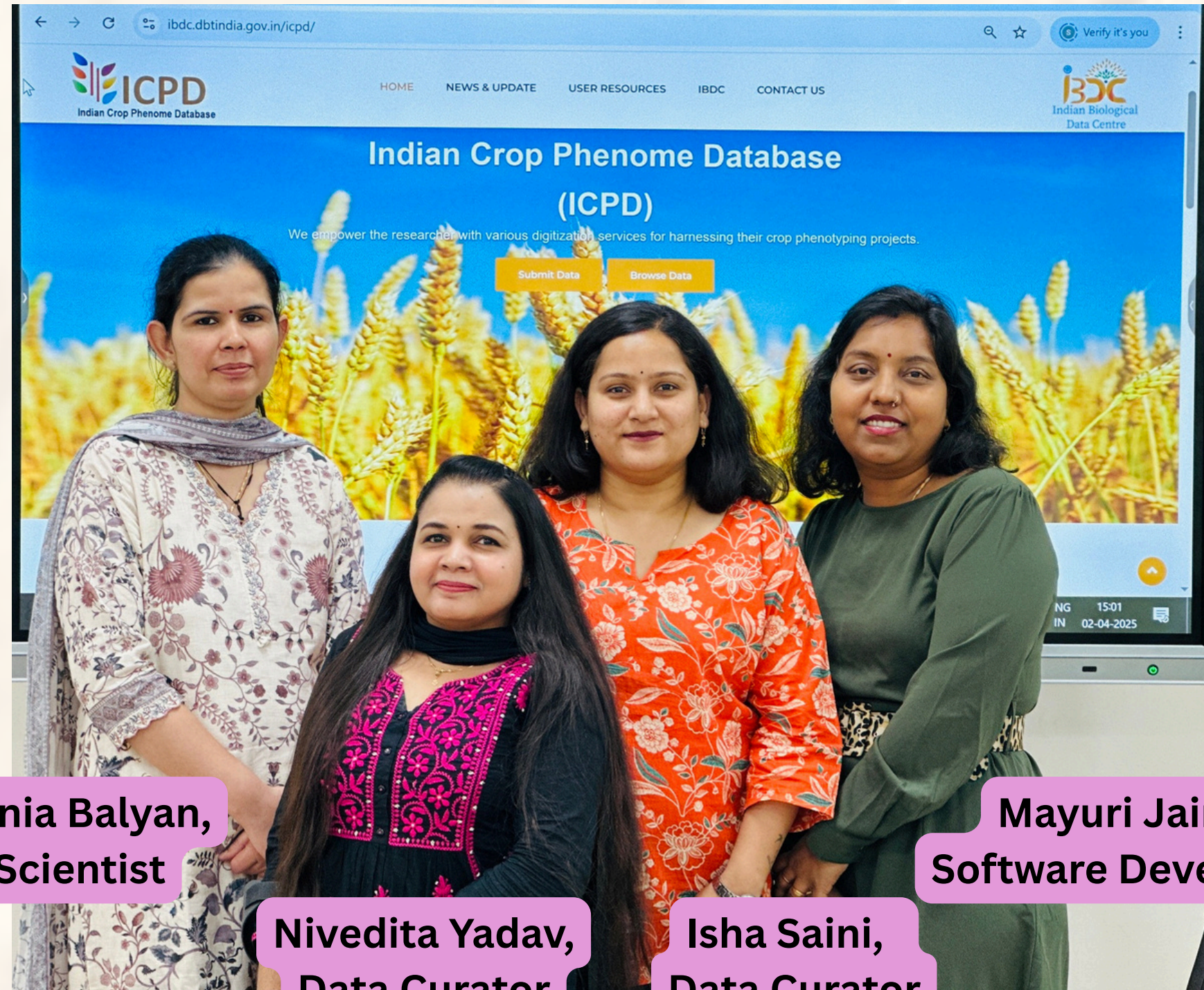






Coming soon.....

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Software Developer, IPR Project

Acknowledgement:

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Pratush Behere (DPMB)
Abhay Shankar Pandey (IBDC)

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National Institute of Immunology
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Dr. Shashank Shekhar,
Head, IBDC



Acknowledgements

- Dr. Rajesh Gokhale (Secretary, DBT)
- Dr. Renu Swarup (Former Secretary, DBT)
- Dr. Arvind Sahu (Executive Director, RCB)
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- Dr. Suchita Ninawe (Advisor, DBT)
- Dr. Saurabh Raghuvanshi (Former Lead, IBDC)
- Dr. Richi Mahajan (Scientist D, DBT)
- Mr. Naveen Kumar (System Administrator, IT, RCB)
- Vipul Adhana (Technical Assistant, IBDC)
- Gautam Kanwal (Technical Assistant, IBDC)
- Locuz



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BIOTECHNOLOGY



International Centre for Genetic
Engineering and Biotechnology



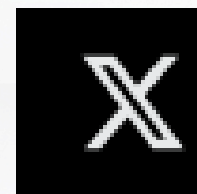
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