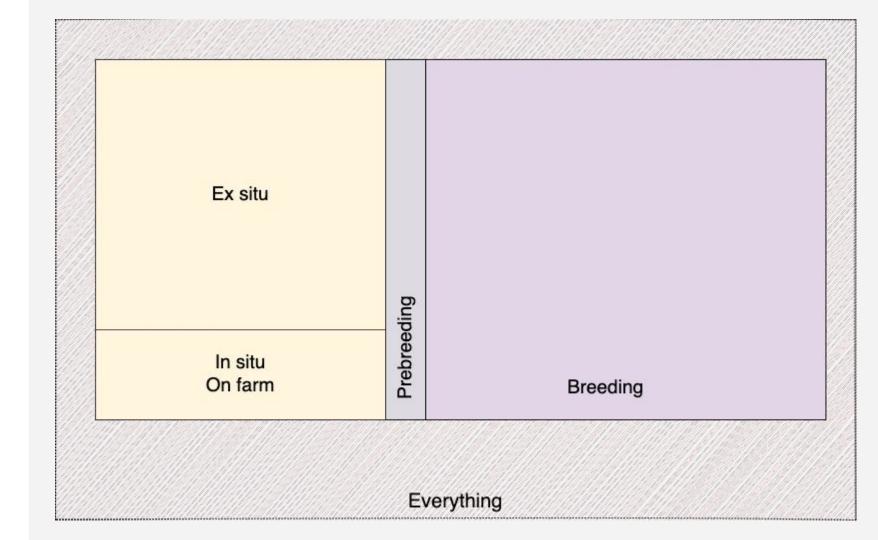
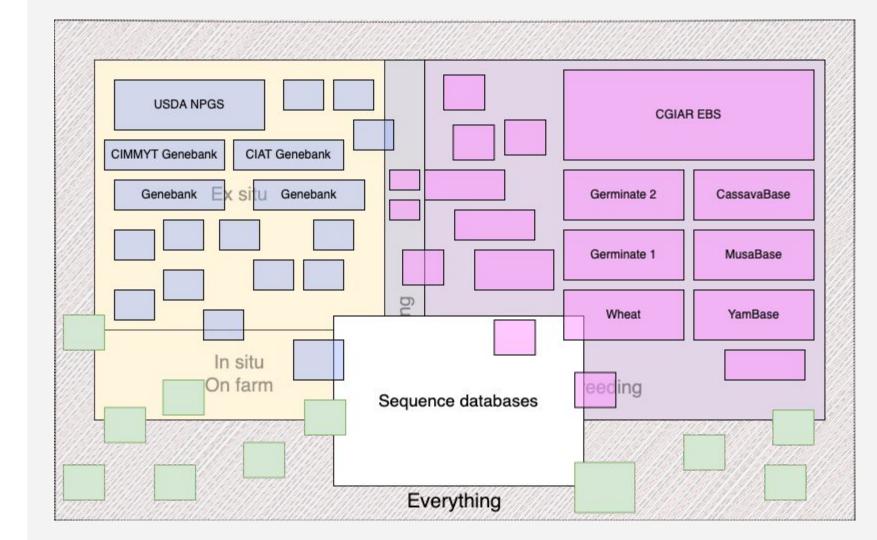
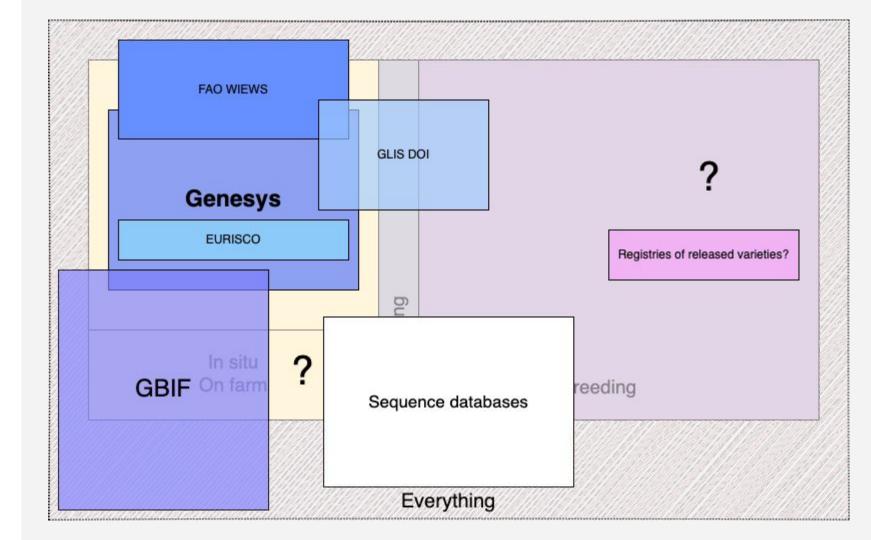


# PGRFA domain







# GGCE / EURISCO / Genesys



## Internal genebank data management

Every genebank needs an information system to effectively maintain their collections.

- Requires a lot of resources
- People **retire**
- Technologies change

Adopt (not adapt) an existing system!

- Learn from the community
- Share with the community



## **Publicly accessible databases**

**Standardization**: Data from different genebanks is made available to users in a standard format.

**Safety** and **security**: Only selected data is exposed to the public. A copy of the data is hosted in a separate database. Master data is behind a firewall.

**Convenience**: More data means better utility for users of genebanks and easier discovery of available material. Economy of scale.

## Genesys

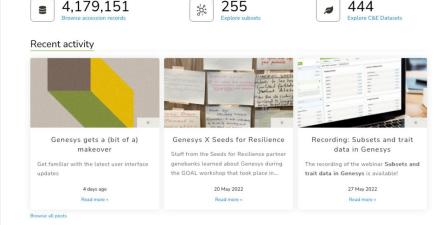
~ Est. 2011 ~

## www.genesys-pgr.org

**Genesys** is an online platform where you can find information about Plant Genetic Resources for Food and Agriculture (**PGRFA**) in crop genebanks worldwide.

**GRIN-Global** and **Genesys** development was initiated in **2008**, in collaboration with USDA and Bioversity, respectively.





Cowpea

Fruit trees

Highlighted crops

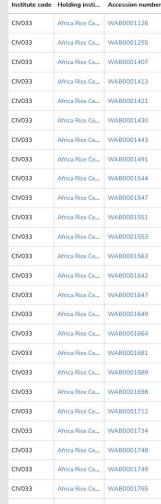


# Scope of Genesys



Genesys focuses on accession-level data about the material in long-term ex situ conservation.

Genesys **does not** host information on *in* situ germplasm, breeding materials or released varieties, unless formally part of genebank collections.



CIV033
CIV033

CIV033	Africa Rice Ce	W
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033	Africa Rice Ce	WAB0001126	TOG 5284
)33	Africa Rice Ce	WAB0001255	TOG 5552
)33	Africa Rice Ce	WAB0001407	TOG 6196
)33	Africa Rice Ce	WAB0001413	TOG 6203
)33	Africa Rice Ce	WAB0001421	TOG 6211
)33	Africa Rice Ce	WAB0001430	TOG 6221
033	Africa Rice Ce	WAB0001443	TOG 6235
)33	Africa Rice Ce	WAB0001491	TOG 6285
033	Africa Rice Ce	WAB0001544	TOG 6339
033	Africa Rice Ce	WAB0001547	TOG 6342
033	Africa Rice Ce	WAB0001551	TOG 6346
033	Africa Rice Ce	WAB0001553	TOG 6349
)33	Africa Rice Ce	WAB0001563	TOG 6359
)33	Africa Rice Ce	WAB0001642	TOG 6441
)33	Africa Rice Ce	WAB0001647	TOG 6449
33	Africa Rice Ce	WAB0001649	TOG 6456
33	Africa Rice Ce	WAB0001664	TOG 6472
033	Africa Rice Ce	WAB0001681	TOG 6489
)33	Africa Rice Ce	WAB0001689	TOG 6497
33	Africa Rice Ce	WAB0001698	TOG 6506
33	Africa Rice Ce	WAB0001712	TOG 6520
033	Africa Rice Ce	WAB0001734	TOG 6542
033	Africa Rice Ce	WAB0001748	TOG 6556
033	Africa Rice Ce	WAB0001749	TOG 6557
)33	Africa Rice Ce	WAB0001765	TOG 6573
)33	Africa Rice Ce	WAB0001781	TOG 6589
033	Africa Rice Ce	WAB0001819	TOG 6629

Accession name	Taxonomy	Crop
TOG 5284	Oryza glaberrima Steud.	Rice
TOG 5552	Oryza glaberrima Steud.	Rice
TOG 6196	Oryza glaberrima Steud.	Rice
TOG 6203	Oryza glaberrima Steud.	Rice
TOG 6211	Oryza glaberrima Steud.	Rice
TOG 6221	Oryza glaberrima Steud.	Rice
TOG 6235	Oryza glaberrima Steud.	Rice
TOG 6285	Oryza glaberrima Steud.	Rice
TOG 6339	Oryza glaberrima Steud.	Rice
TOG 6342	Oryza glaberrima Steud.	Rice
TOG 6346	Oryza glaberrima Steud.	Rice
TOG 6349	Oryza glaberrima Steud.	Rice
TOG 6359	Oryza glaberrima Steud.	Rice
TOG 6441	Oryza glaberrima Steud.	Rice
TOG 6449	Oryza glaberrima Steud.	Rice
TOG 6456	Oryza glaberrima Steud.	Rice
TOG 6472	Oryza glaberrima Steud.	Rice
TOG 6489	Oryza glaberrima Steud.	Rice
TOG 6497	Oryza glaberrima Steud.	Rice
TOG 6506	Oryza glaberrima Steud.	Rice
TOG 6520	Oryza glaberrima Steud.	Rice
TOG 6542	Oryza glaberrima Steud.	Rice
TOG 6556	Oryza glaberrima Steud.	Rice

Oryza glaberrima Steud.	Rice	
Oryza glaberrima Steud.	Rice	1
Oryza glaberrima Steud.	Rice	1
Oryza glaberrima Steud.	Rice	-
Oryza glaberrima Steud.	Rice	1
Oryza glaberrima Steud.	Rice	
Oryza glaberrima Steud.	Rice	1
Oryza glaberrima Steud.	Rice	1
Oryza glaberrima Steud.	Rice	1
Oryza glaberrima Steud.	Rice	1
Oryza glaberrima Steud.	Rice	1
Oryza glaberrima Steud.	Rice	1
Oryza glaberrima Steud.	Rice	1
Oryza glaberrima Steud.	Rice	1
Oryza glaberrima Steud.	Rice	1
Oryza glaberrima Steud.	Rice	1
Oryza glaberrima Steud.	Rice	1
Oryza glaberrima Steud.	Rice	1
Oryza glaberrima Steud.	Rice	1
Oryza glaberrima Steud.	Rice	1

Oryza glaberrima Steud. Rice

Crop name Crop

Rice

Biological status of accession Provena

Nigeria

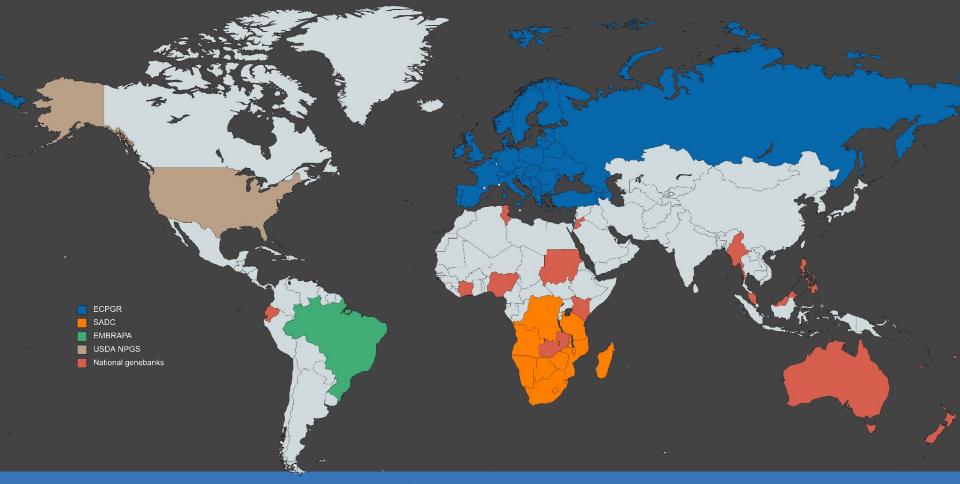
Traditional cultivar/Landrace

Traditional cultivar/Landrace

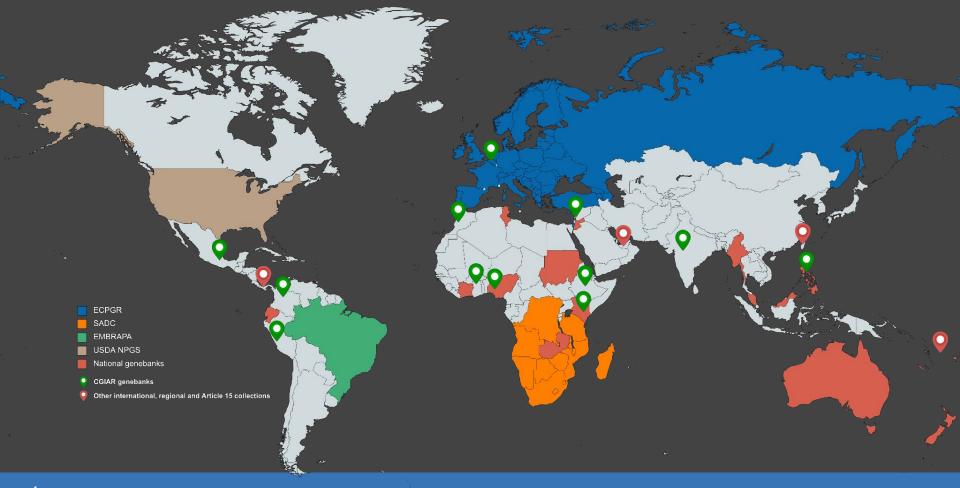
ud.	Rice	Rice	Traditional cultivar/Landrace	Guinea
ud.	Rice	Rice	Traditional cultivar/Landrace	Guinea
ud.	Rice	Rice	Traditional cultivar/Landrace	Nigeria
ud.	Rice	Rice	Traditional cultivar/Landrace	Nigeria
ud.	Rice	Rice	Traditional cultivar/Landrace	Liberia
ud.	Rice	Rice	Traditional cultivar/Landrace	Liberia
ud.	Rice	Rice	Traditional cultivar/Landrace	Liberia
ud.	Rice	Rice	Traditional cultivar/Landrace	Liberia
ud.	Rice	Rice	Traditional cultivar/Landrace	Liberia
ud.	Rice	Rice	Traditional cultivar/Landrace	Liberia
ud.	Rice	Rice	Traditional cultivar/Landrace	Liberia
ud.	Rice	Rice	Traditional cultivar/Landrace	Mali
ud.	Rice	Rice	Traditional cultivar/Landrace	Mali
ud.	Rice	Rice	Traditional cultivar/Landrace	Mali
ud.	Rice	Rice	Traditional cultivar/Landrace	Liberia
ud.	Rice	Rice	Traditional cultivar/Landrace	Liberia
ud.	Rice	Rice	Traditional cultivar/Landrace	Liberia
ud.	Rice	Rice	Traditional cultivar/Landrace	Liberia
ud.	Rice	Rice	Traditional cultivar/Landrace	Liberia
ud.	Rice	Rice	Traditional cultivar/Landrace	Liberia
ud.	Rice	Rice	Traditional cultivar/Landrace	Liberia
ud.	Rice	Rice	Traditional cultivar/Landrace	Liberia
ud.	Rice	Rice	Traditional cultivar/Landrace	Liberia
ud.	Rice	Rice	Traditional cultivar/Landrace	Liberia

Traditional cultivar/Landrace











# **Tools for** genebanks



# https://validator.genesys-pgr.org

Spelling of taxonomic names based on GRIN

Taxonomy

GIS test of coordinates, country of provenance



**GENUS** 

Sorghum

Sorghum

Sorghum

Sorghum

Sorghum

Sorgho

Sorghum

Sorghum

GENUS\_check

OK

OK

OK

OK

OK

OK

OK

Sorghum; Sorgum

35092 true true

true

true

true

true

true

GRINTAX\_species

35092 35092 true

GRINTAX\_speciesId

	bicolor			
******	ок	35092	true	uningnoon
	<i>(</i> 2 ) )			

SPECIES\_check

OK

OK

OK

SPECIES

bicolor

bicolor

bicolor

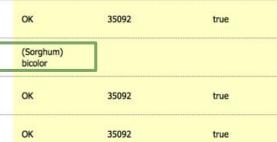
bicolo

bicolor

bicolor

bicolor

bicolor



35092

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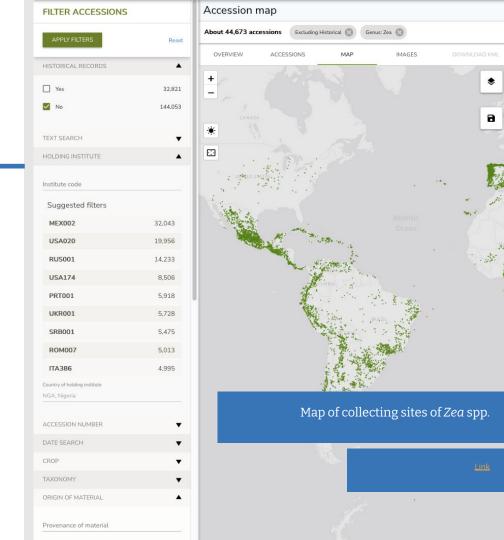


# Exploring PGRFA data



- Full-text search and filtering
- Maps
- Data summaries (Overviews)
- Finding accessions by climate
- Finding similar/duplicate accessions
- History of updates to passport data
- Initiating requests for material
- Coming soon: Subsetting Tool





## Genesys and genebank operations





## **Acquisition (improving collections):**

- Which material is already conserved?
- What is unique? Gap analysis
- Any updates relevant to material received from other genebanks?

## **Management (updating Genesys):**

- Availability
- Type of germplasm maintenance

## **Distribution (for use):**

Finding and requesting material



# Beyond passport data



- Accession **images** and scanned collecting forms
- Collection subsets (core- and mini-core collections, reference sets, etc.)
- Datasets of trait observations annotated with trait descriptors and other metadata



### Aegilops tauschii mini-core collection

The Wheat Genetics Resource Center at Kansas State University has genotyped a collection of 600 Ae. tauschii accessions and selected a set of 40 lines that are the most genetically diverse and capturing more than 95% of the allelic diversity. This mini-core includes 28 accessions from Lineage 1 [strangulata type] and 12 from Lineage 2 [eusquarrosa type]. The mini-core represents 13 countries spanning the distribution of Ae. tauschii, three accession are of unknown origin; 27.5% are from Iran and 17.5% are from Afghanistan, countries with the largest density of this species.

Single nucleotide polymorphism (SNP) discovery and genotyping was performed in single step with Tassel 5 GBSv2 pipeline using the *Ae. tauschii* genome assembly (Aet v4.0; NCBI BioProject

PRJNA341983) as the reference. Tassel was run with bowtie2 aligner for tags mapping in Linux HPC environment via shell script. Population level SNP filtering was performed and SNPs with a minor allele frequency (MAF) less than 0.01 and missing data more than 20% were removed. SNPs with heterozygosity greater than 5% were removed because Ae. tauschii accessions are highly inbred. Individual samples with more than 80% missing SNP calls and more than 5% heterozygosity were also removed.

All SNPs were used to select a representative core-set from the Ae. tauschii collection. The core-set was selected in two steps. First, the software package PowerCore was used with default settings, which selects the lines to retain most diverse alleles by

implementing advanced M (maximization) strategy. Then the number of selected accessions was further reduced with a phenotypically guided selection using the available phenotypic data for a leaf rust composite culture, stem rust race TTKSK, and Hessian fly biotype D resistance. The diversity captured by the MiniCore was assessed by the percent segregating SNPs present in the selected accessions relative to the whole collection.

#### Reference:

Genomic analysis confirms population structure and identifies inter-lineage hybrids in *Aegilops tauschii*. 2019. Singh N, Wu S, Tiwari V, Sehgal S, Raupp J, Wilson D, Abbasov M, Gill B, and Poland J. Frontiers in Plant Science. https://doi.org/10.3389/fpls.2019.00009

Type of subset	Selective
Method of selection	Genoyping-by-sequencing
Crop	Wheat
Number of accessions	40
Institute	Wheat Genetics Resource Center
Creation date	2015
Source	☑ https://doi.org/10.3389/fpls.2019.00009

## Subset metadata

#### Subset creators

Data curator	Eduard A
Data curator	Bikram S
Data curator	Jesse Po

Genomic analysis confirms population structure and identifies inter-lineage hybrids in *Aegilops tauschii*. 2019. Singh N, Wu S, Tiwari V, Sehgal S, Raupp J, Wilson D, Abbasov M, Gill B, and Poland J. Frontiers in Plant Science.

Data manager W. John Raupp Kansas State University

Data curator Sunish Sehgal South Dakota State University [current affiliation]

# Public genebank websites

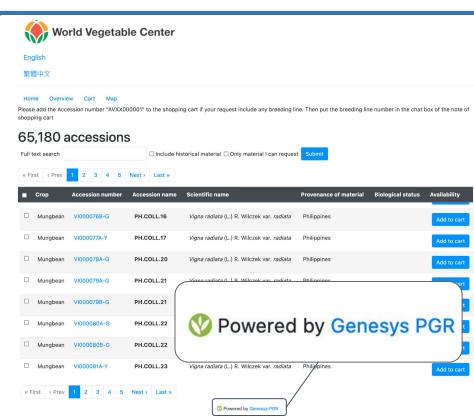


## **Embedded Genesys**

Instead of hosting and maintaining a separate database, paying for a web server and setting up a new domain name, genebanks can use Embedded Genesys to integrate their data from Genesys directly into existing institutional websites with a few lines of Javascript.

The latest feature added to Embedded Genesys is the Similarity Search. When an accession is marked as unavailable, users can search for similar accessions and request one of the provided alternatives instead.





# Summary



## Genebank + GGCE

- Manage all data related to PGRFA in your collection and effectively operate your ged
- Community knowledge exchange, support, technical solutions

## **ECPGR + Eurisco**

- Regional
- Knowledge exchange, standards, support, technical solutions

## Genesys

- Cross-regional and global
- Knowledge exchange, standards, support, technical solutions



## www.genesys-pgr.org



What's new?

https://www.genesys-pgr.org/content/news

Webinar: How to use Genesys?

https://www.genesys-pgr.org/content/news/129/recording-how-to-use-genesys

Webinar: Subsets and trait data in Genesys

https://www.genesys-pgr.org/content/news/136/recording-subsets-and-trait-data-in-ge

<u>nesys</u>

Searching for similar accessions

https://www.genesys-pgr.org/content/news/104/searching-for-similar-accessions

Get in touch!

helpdesk@genesys-pgr.org





www.croptrust.org