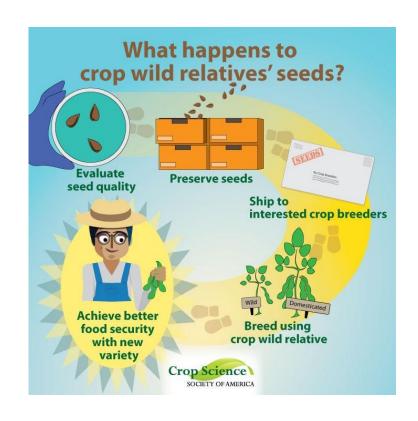


# FIGS: Combining phenotypic, environmental and genomic data to mine collections

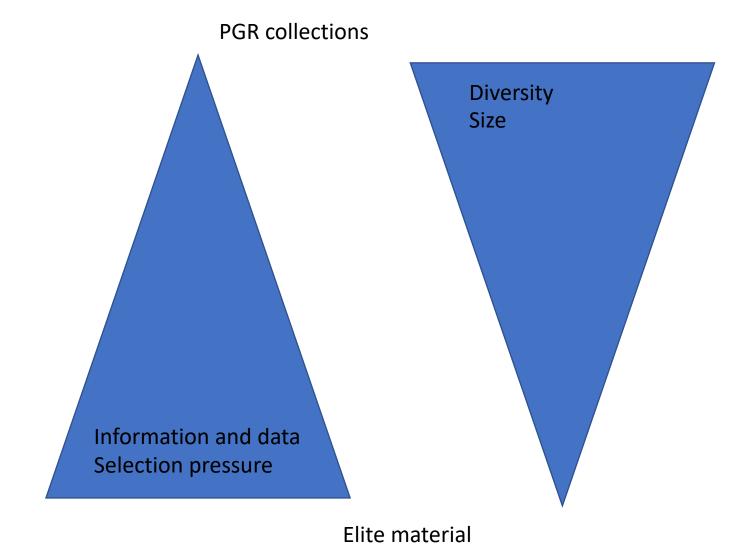


Zakaria **Kehel** z.kehel@cgiar.org





## Background

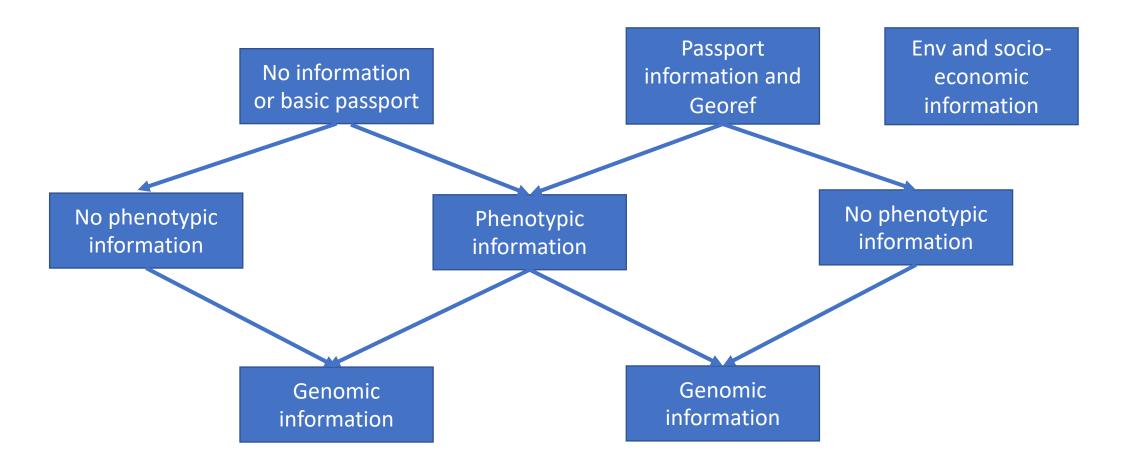


## Digital genetic resources

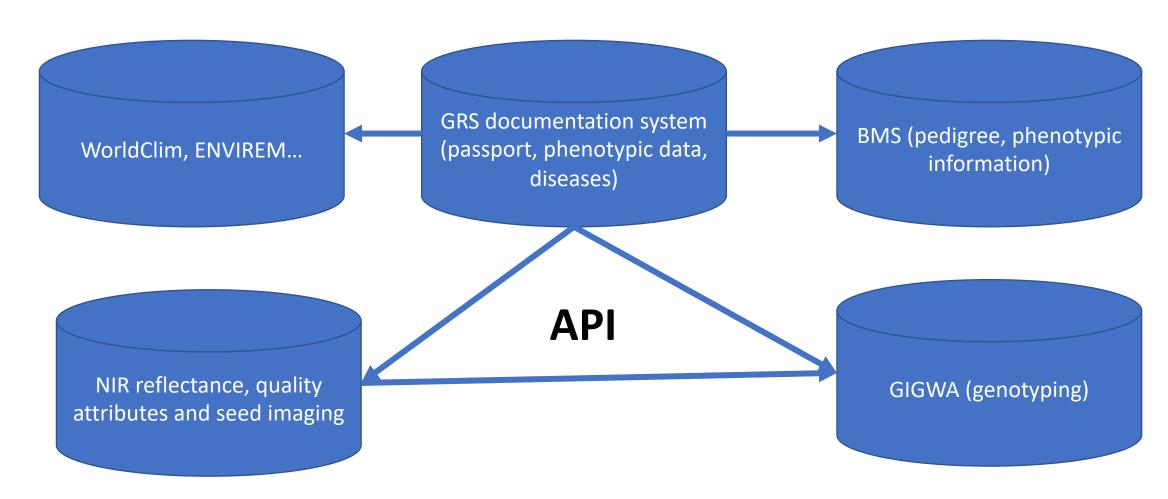




#### What to do now?



## Data availability



#### **Focused Identification of Germplasm Strategy (FIGS)**

#### **FIGS Approach**

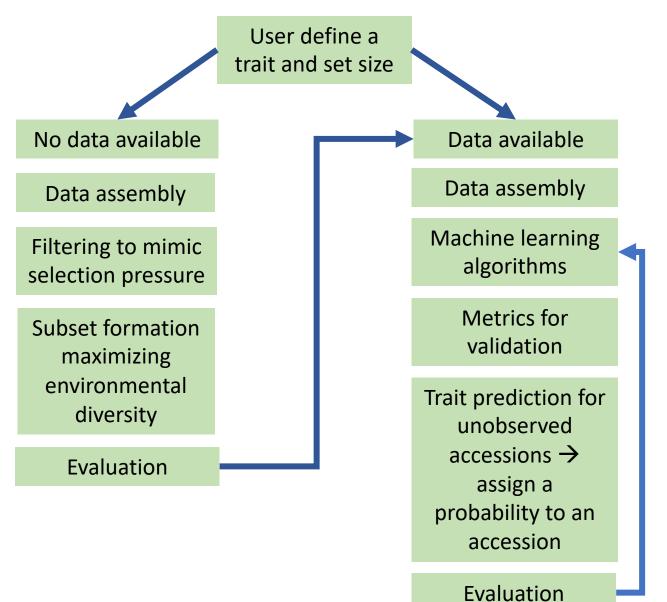


By **linking** traits, environments (and associated selection pressures) with genebank accessions (e.g. landraces and crop relatives) we can **focus** in on those accessions most likely to possess **trait specific genetic variation**.

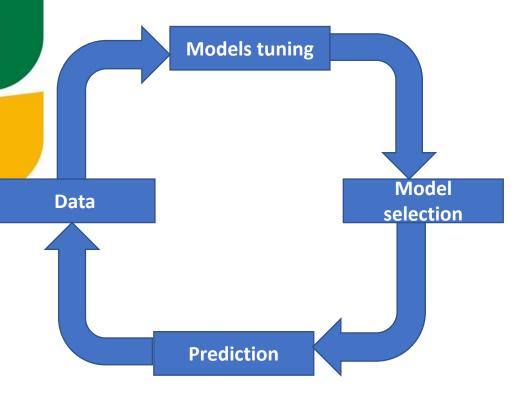
Two methods resulting from the FIGS approach:

- 1. Filtering method
- 2. 'Predictive' modeling method

## FIGS pathways



# Durum wheat Grain filling period example

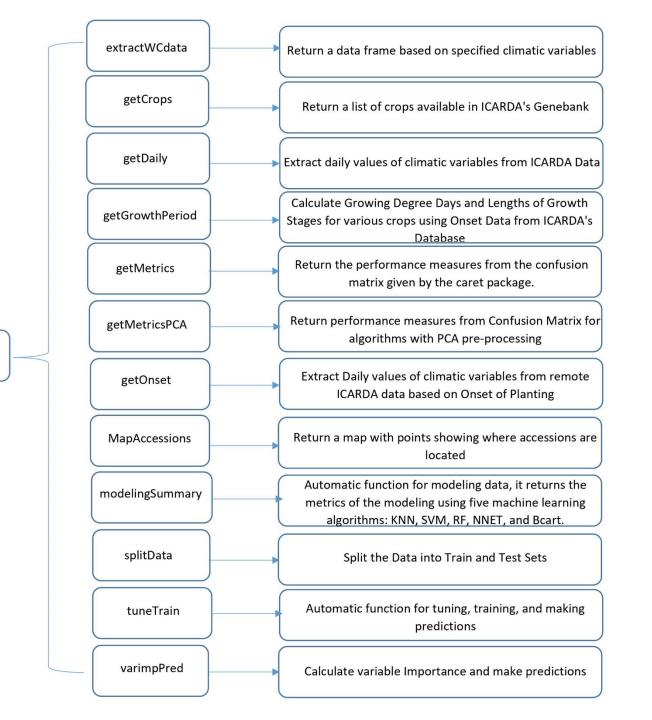


FIGS modeling pathway

#### **Predictive characterization**

Performance Measures		k-Nearest Neighbours	Random Forest	Support Vector Machine	
	Accuracy	0.834	0.838	0.817	
95% CI		(0.799, 0.865)	(0.804, 0.868)	(0.781, 0.849)	
No Information Rate		0.762	0.762	0.762	
P-Value [Acc > NIR]		3.58E-05	1.37E-05	0.001423371	
Kappa		0.563	0.557	0.467	
Sensitivity		0.722	0.675	0.54	
Specificity		0.869	0.889	0.903	
True Positive		91	85	68	
True Negative		351	359	365	
False Positive		53	45	39	
False Negative		35	41	58	

High accuracy showing that there is a strong relationship between GFP and longterm climatic conditions



icardaFIGSr

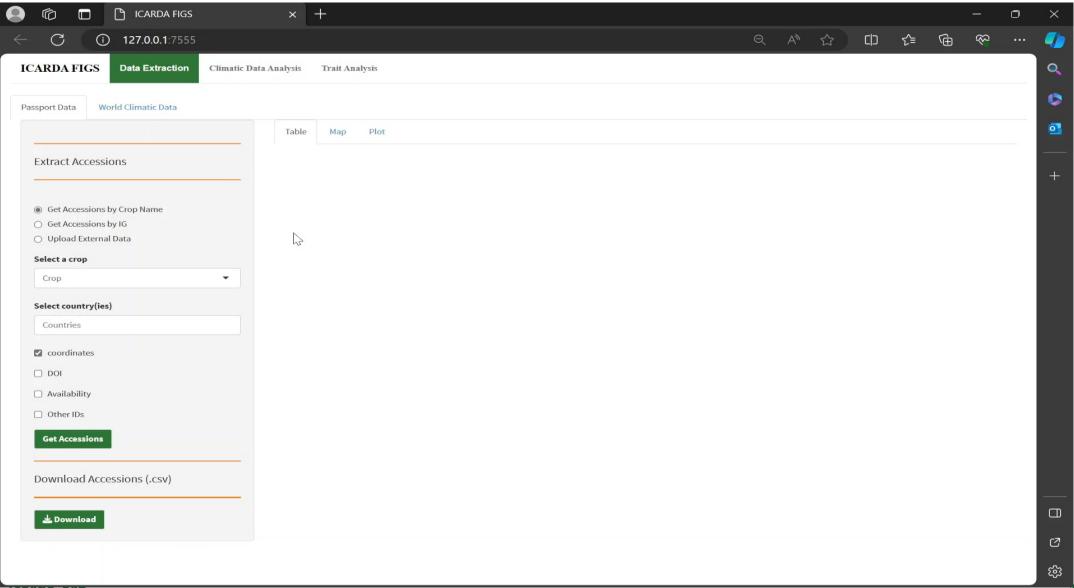
## **Tools:** R package

#### TuneTrain Splitting the data If parallel computing is True, then do parallel computing Choose a method RF SVM **NNET Bcart** KNN Tuning Training Choose the best parameter for training **Training** Target variable is factor or numeric? Factor Numeric Class probabilities are calculated for Predicted values are calculated each class A plot of residuals versus predicted A ROC curve is created values is created Return a list containing a model object, predictions, class probabilities, and the training and test data sets

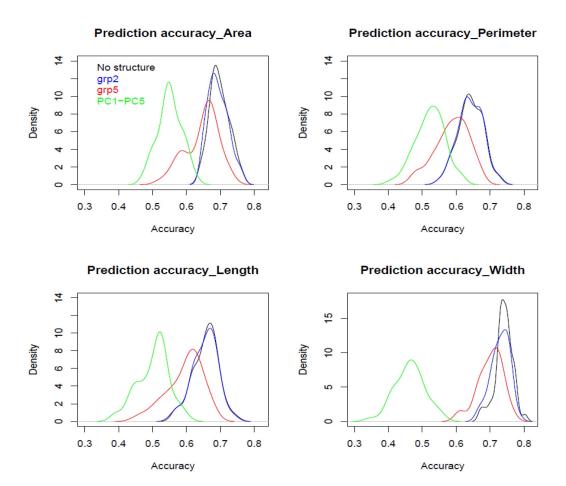
### **Tools:** R package

- TuneTrain is an automatic function for tuning and training data, it returns a list containing a model object, predictions, class probabilities, and the training and test data sets
- TuneTrain(data, y, p = 0.7, method = method, parallelComputing = F, length = 10, control = "repeatedcv", number = 10, repeats = 10, process = c('center', 'scale'), summary= multiClassSummary,positive, ...)

### **Tools: GUI**



## Adding value Genetic resources: FIGS augmented with Genomic selection for wheat quality example

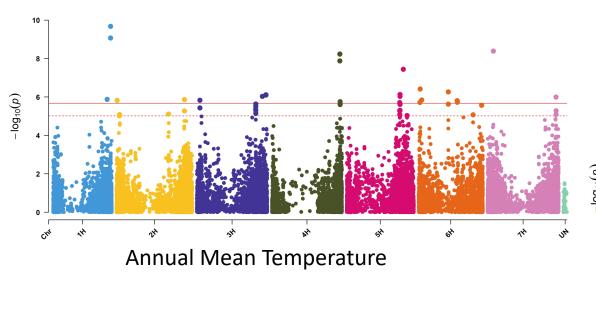


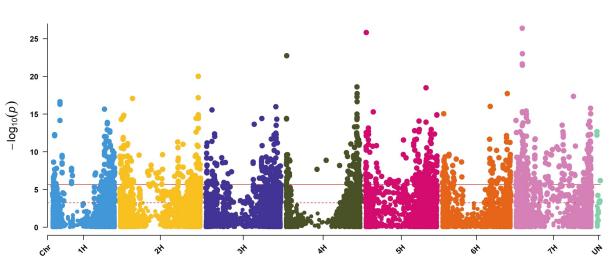
		Area	Perimeter	Length	Width	ColCha1	ColCha2	ColCha3
All	Mean	0.70	0.64	0.66	0.74	0.69	0.71	0.66
	Maximum	0.76	0.73	0.74	0.80	0.78	0.80	0.73
	Minimum	0.64	0.55	0.57	0.67	0.60	0.62	0.59
All-Env	Mean	0.75	0.70	0.70	0.78	0.75	0.76	0.72
	Maximum	0.78	0.75	0.77	0.81	0.79	0.80	0.75
	Minimum	0.66	0.59	0.59	0.69	0.61	0.63	0.60
Env	Mean	0.06	0.04	0.03	0.10	0.03	0.05	0.04
	Maximum	0.20	0.19	0.22	0.29	0.20	0.25	0.24
	Minimum	-0.12	-0.19	-0.16	-0.07	-0.16	-0.13	-0.13
grp1	Mean	0.73	0.76	0.79	0.65	0.58	0.63	0.58
	Maximum	0.82	0.86	0.88	0.75	0.78	0.80	0.71
	Minimum	0.64	0.67	0.69	0.49	0.42	0.47	0.40
grp2	Mean	0.67	0.54	0.51	0.73	0.73	0.74	0.69
	Maximum	0.74	0.67	0.65	0.80	0.79	0.79	0.75
	Minimum	0.58	0.46	0.40	0.65	0.63	0.64	0.61

- GS shows efficiency in predicting seed morphology in Genebank wheat collection
- Combining FIGS and GS showed better predictability
- Increase knowledge on quality attributes which was bottleneck in the past



### Adding value to PGR: eWAS & GWAS



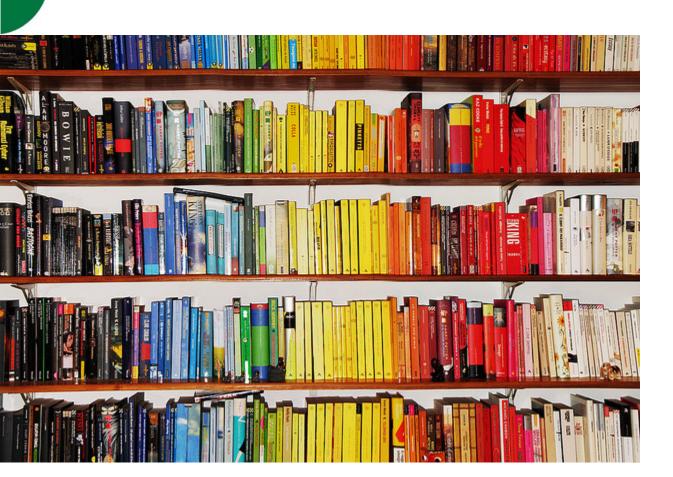


- eWAS revealed several regions for adaptation to Temp, Rain and soil
- Understand adaptation of CRW to different environments
- Increase knowledge to identify sources of adaptation to abiotic stress from PGR

**Annual Precipitation** 

#### Almost there? ©

- Organizing, curating and connecting different datasets
- Take advantage of data mining and machine learning technologies
- Increasing predictability with what is available







Thank you! Questions/Comments?



