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## Norms of reaction for maritime pine

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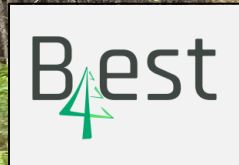
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*B4EST international conference – Managing Forest Genetic Resources for an uncertain future*

# **Norms of reaction for maritime pine**

***Session A** : Accelerating breeding to cope with new challenges and uncertain future – 20<sup>th</sup> June 2022*

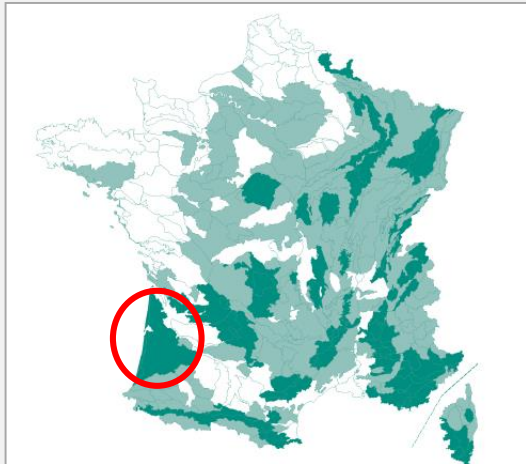
Victor Papin, INRAE (UMR BIOGECO Pierroton)  
Laurent Bouffier, INRAE (UMR BIOGECO Pierroton)  
Léopoldo Sanchez, INRAE (UMR BioForA Orléans)



# Maritime pine (*Pinus Pinaster*) breeding in France

## « Landes de Gascogne » forest

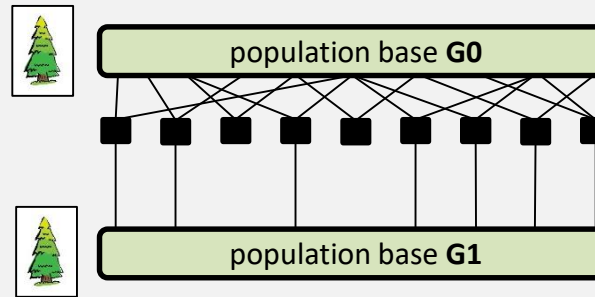
- 0,8 million hectares (24% of French wood harvest)
- Main uses in carpentry, joinery, stationery



- Plantations with **improved seedlings**
- Mainly based on the **Landes provenance**

## Recurrent selection scheme

- Started in the ~1960 → 3rd generation today

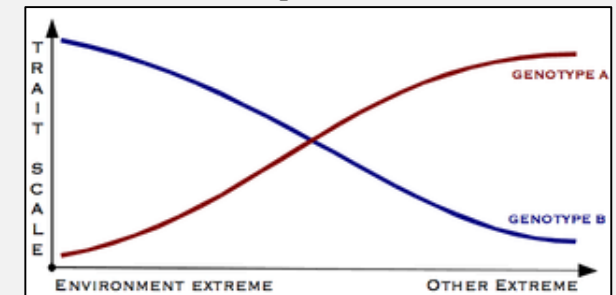


- Targeted traits :
  - **Growth** (height, circumference, straightness)
  - **Adaptation** to the environment
  - **Wood quality**
- **Genetic values** estimated with **pedigree**

## Dynamic phenotypes : NoR

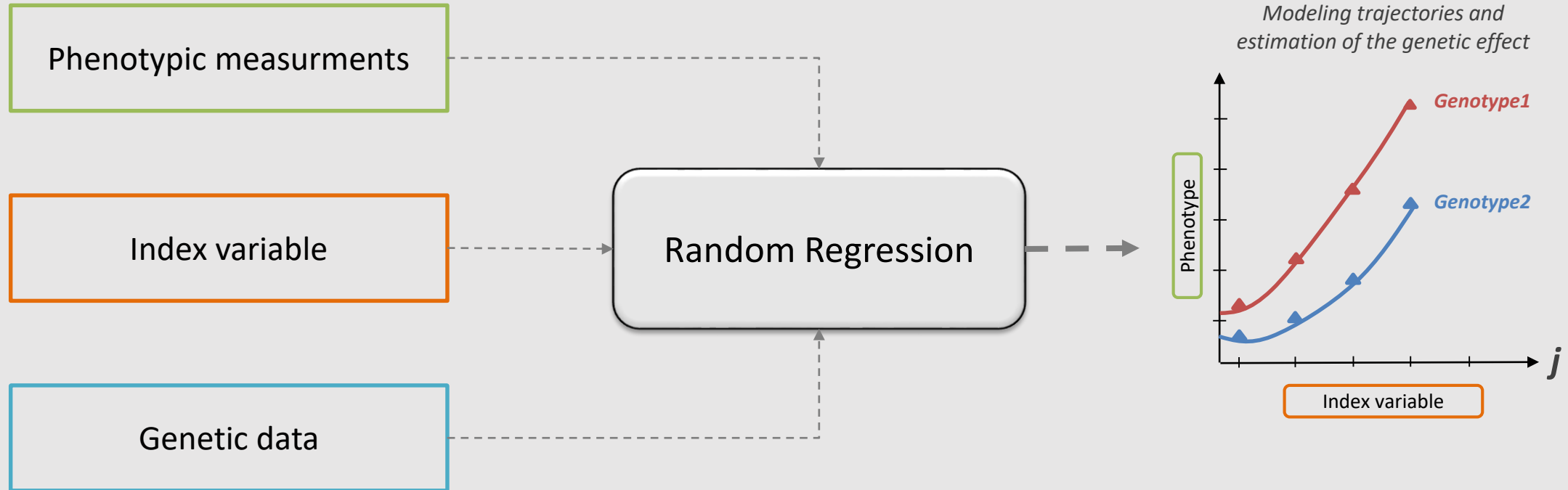
- Classical selection focuses on **final growth traits**
- Integrative phenotypes with **no consideration of the environment**

NoR : pattern of phenotypic expression of a single genotype across a range of environments

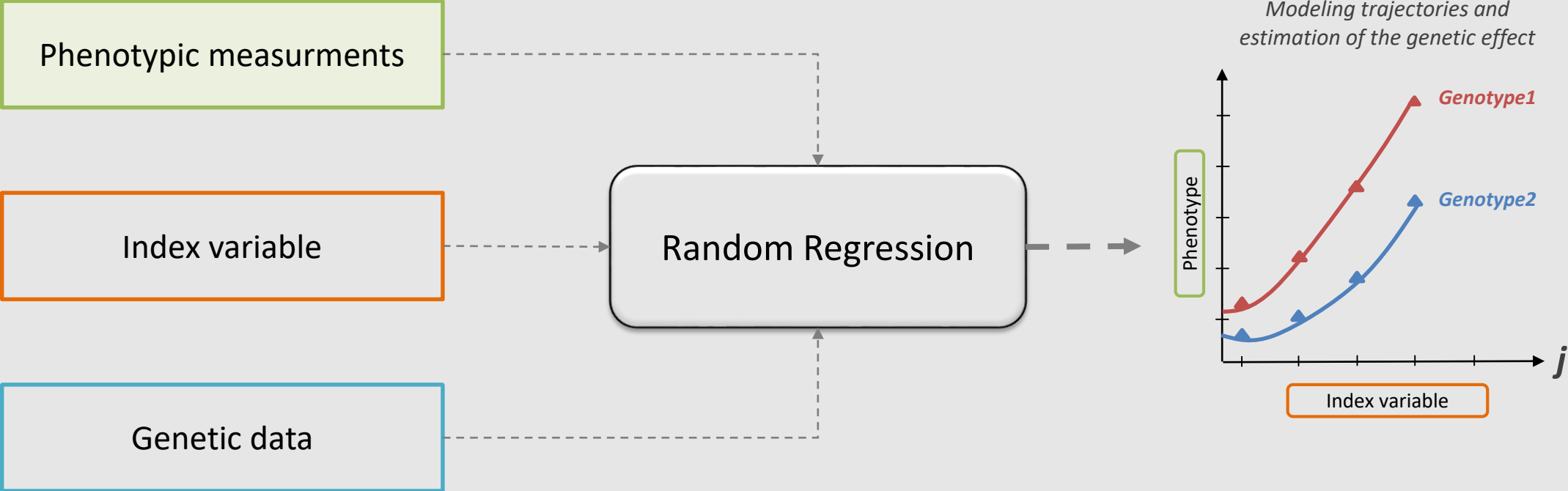


- Consideration of **dynamic** and **explicative** traits
- Evaluation and prediction of **genetic values** taking account of the **environnement** (even in unobserved environments)

# Construction of norms of reaction

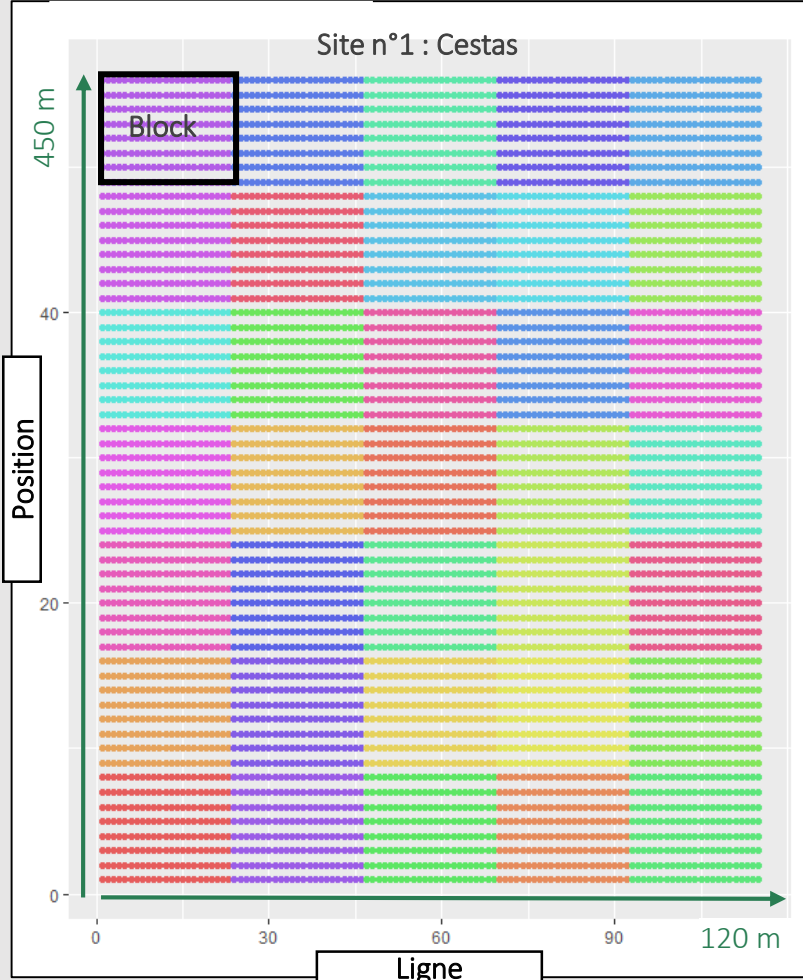


# Construction of norms of reaction



# Experimental design and measurements

Experimental design for maritime pine (6300 trees)

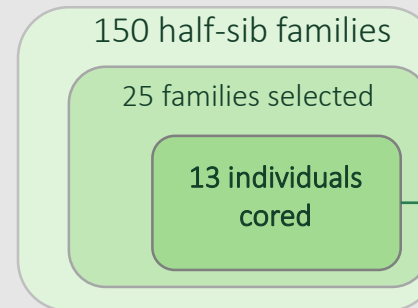


## Experimental design

- 2 locations : Cestas (humid) & Escource (dry)
- Installation in **1996** : **26** years old trees
- **150** half-sib families with **35** individuals/family → **6300** trees per site
- **Complete** block design (1 individual of each family per block)

## Phenotypic measurements :

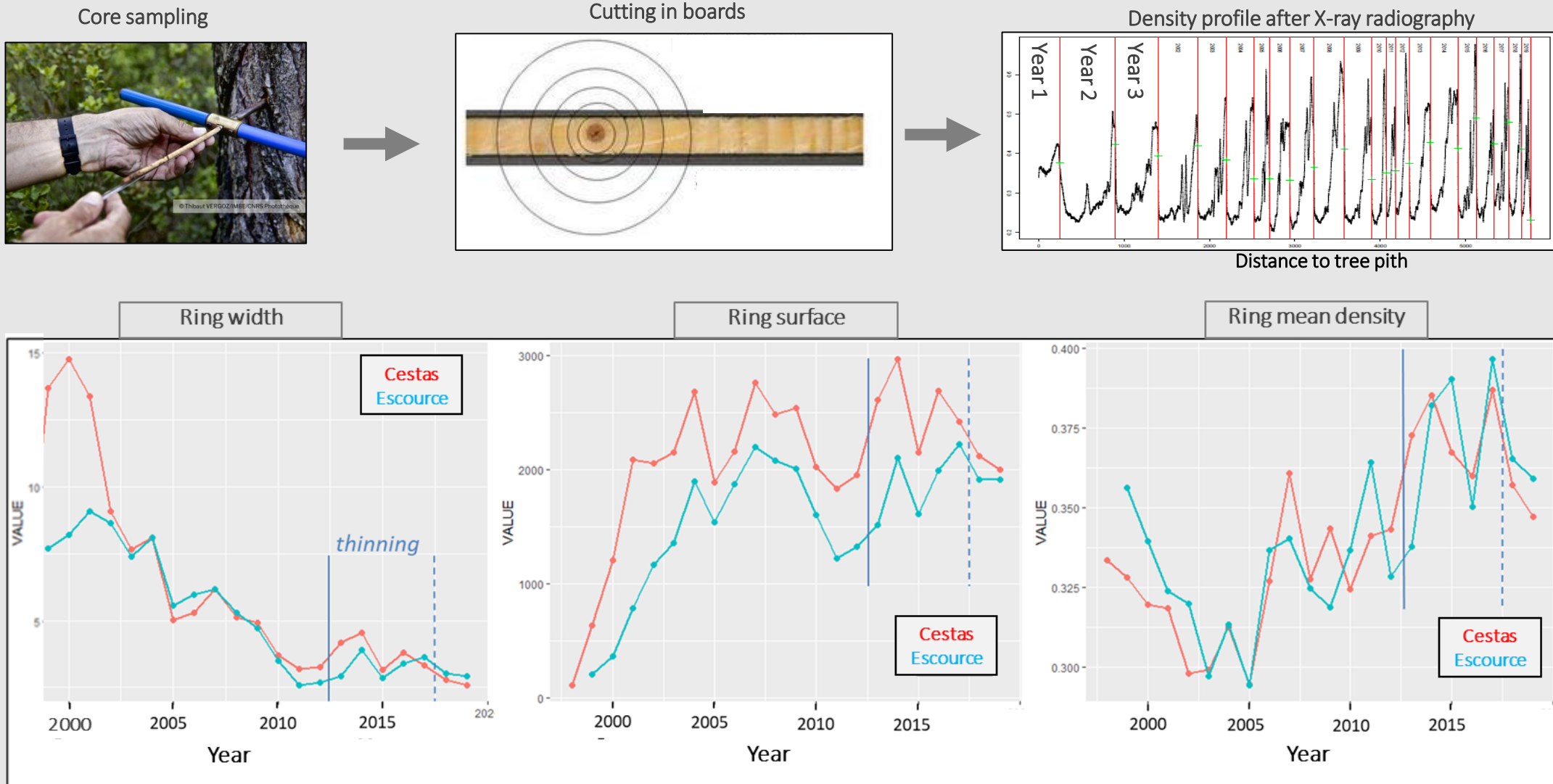
- Classical growth measures at different ages (height, circumference, straightness)
- Core sampling of **325** trees per site :



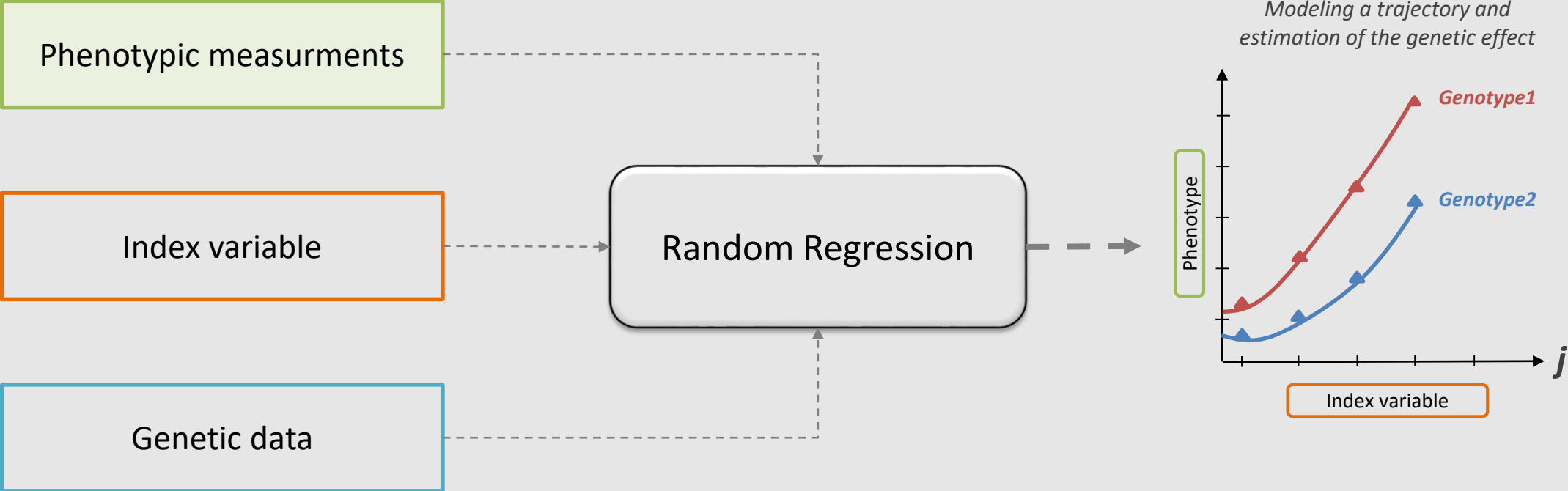
## Environmental measurements :

- Conventional annual climate measurements (temperature, rainfall...)

# Experimental design and measurements

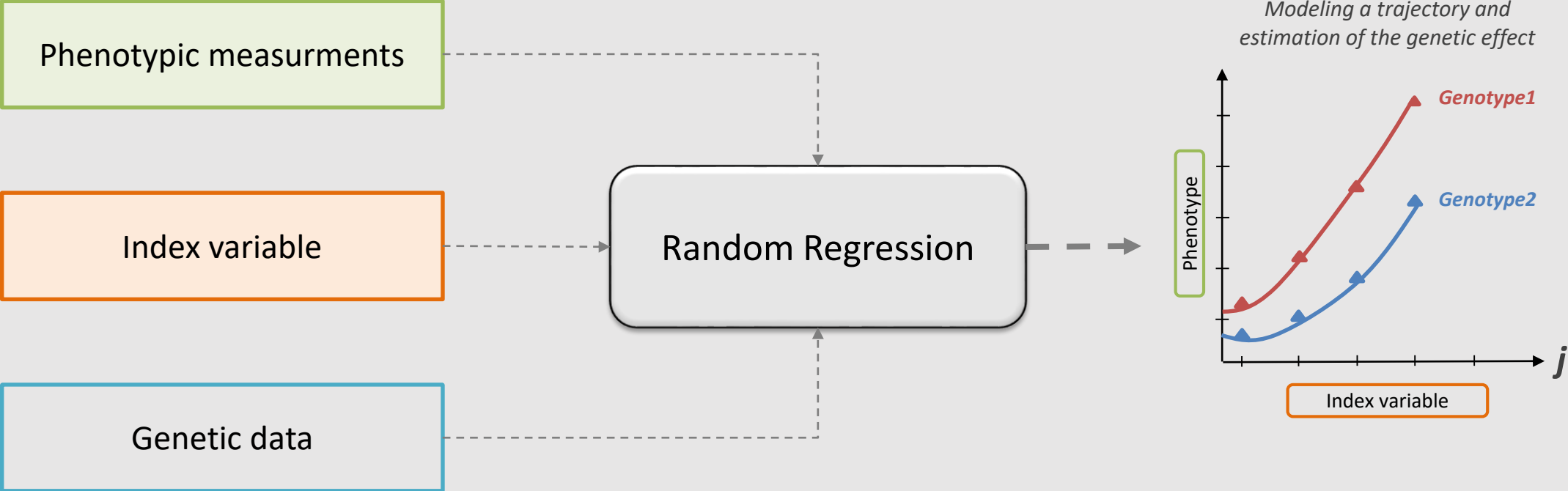


# Construction of norms of reaction

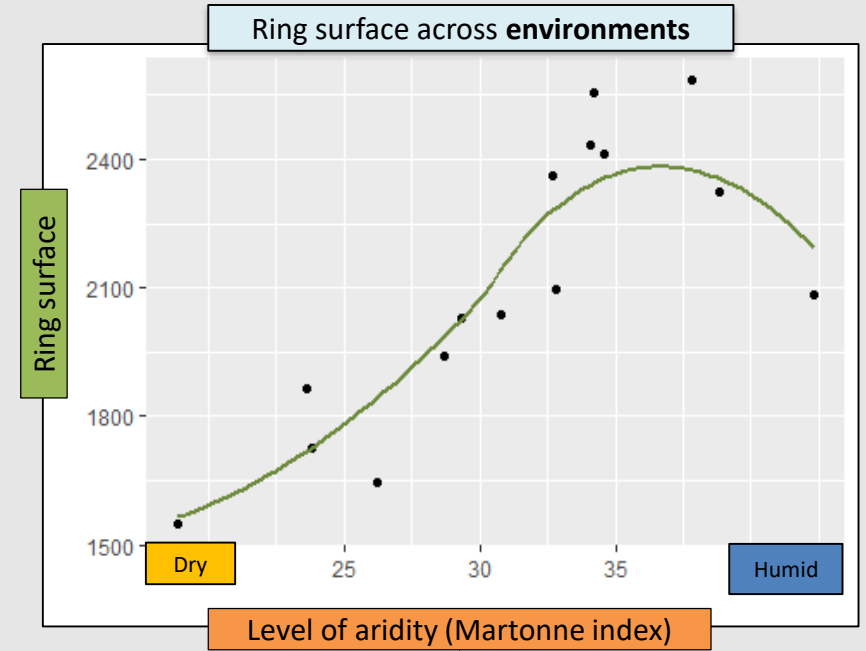
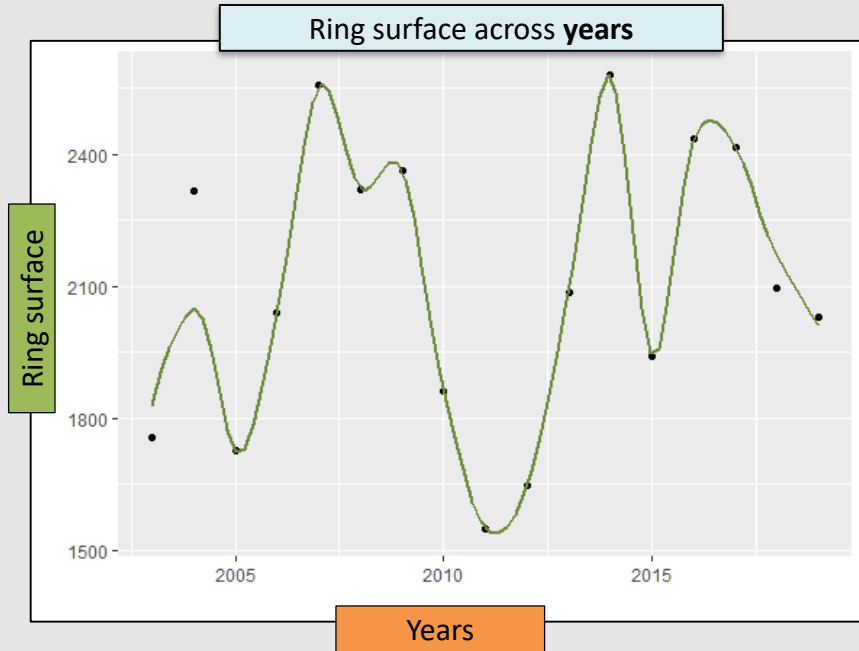




# Construction of norms of reaction



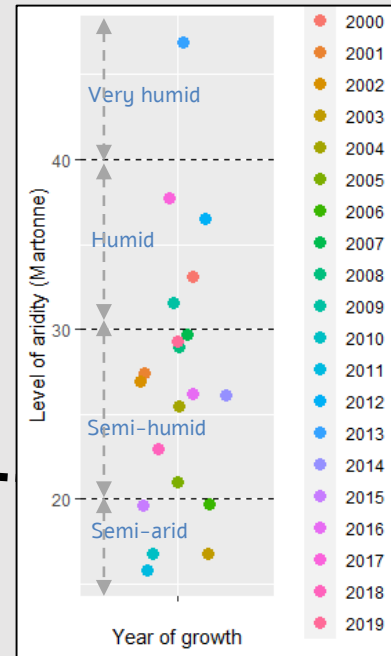
# Construction of norms of reaction



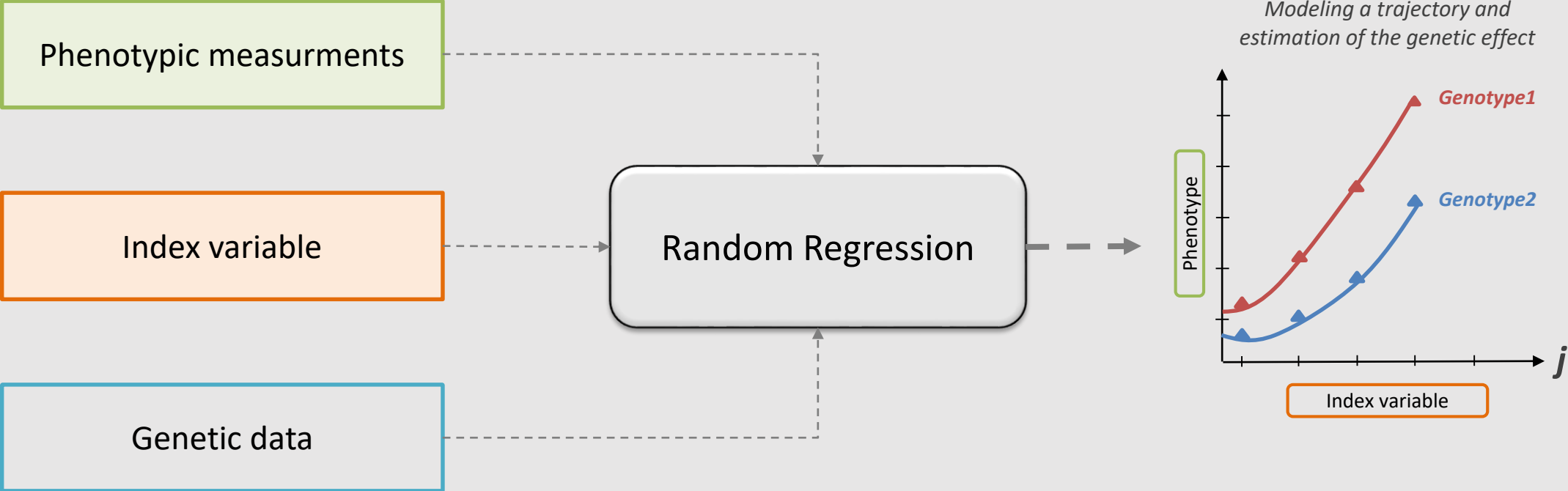
For each year, characterization of global aridity with Martonne index

$$I = \frac{12p}{t + 10}$$

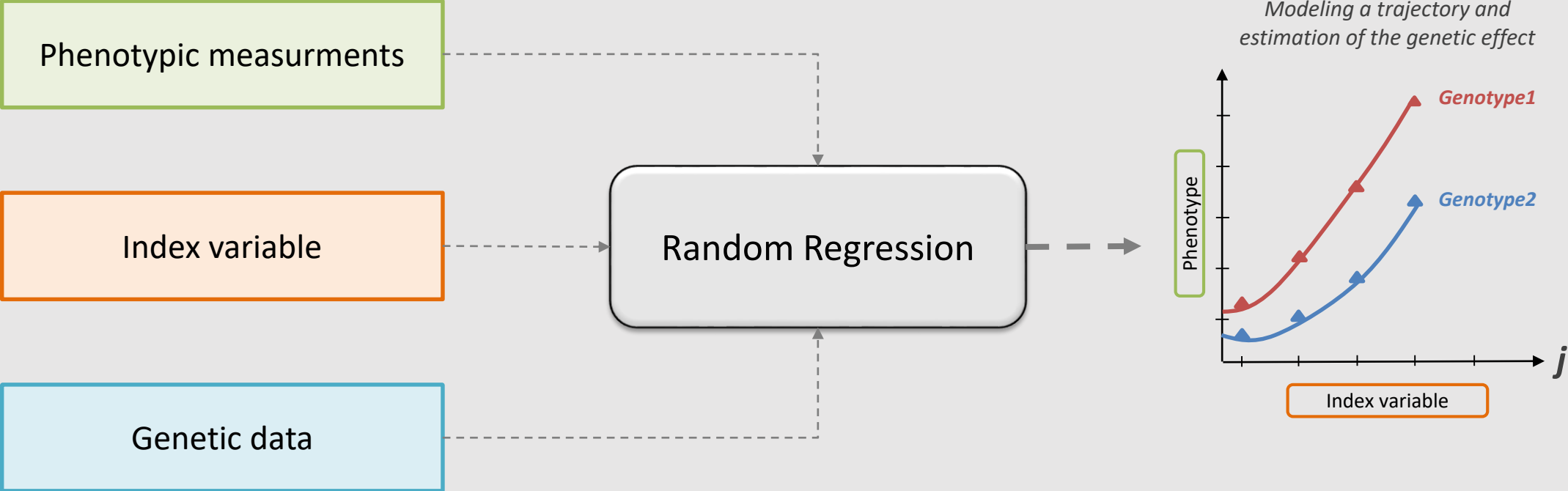
$p$  : cumulative rainfall (mm)  
 $t$  : mean temperature (°C)



# Construction of norms of reaction

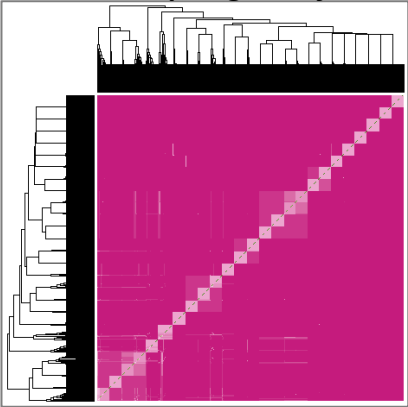


# Construction of norms of reaction

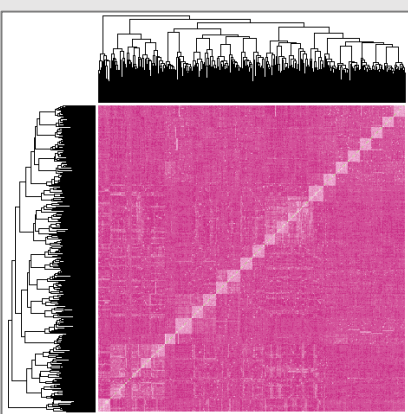


# Construction of norms of reaction

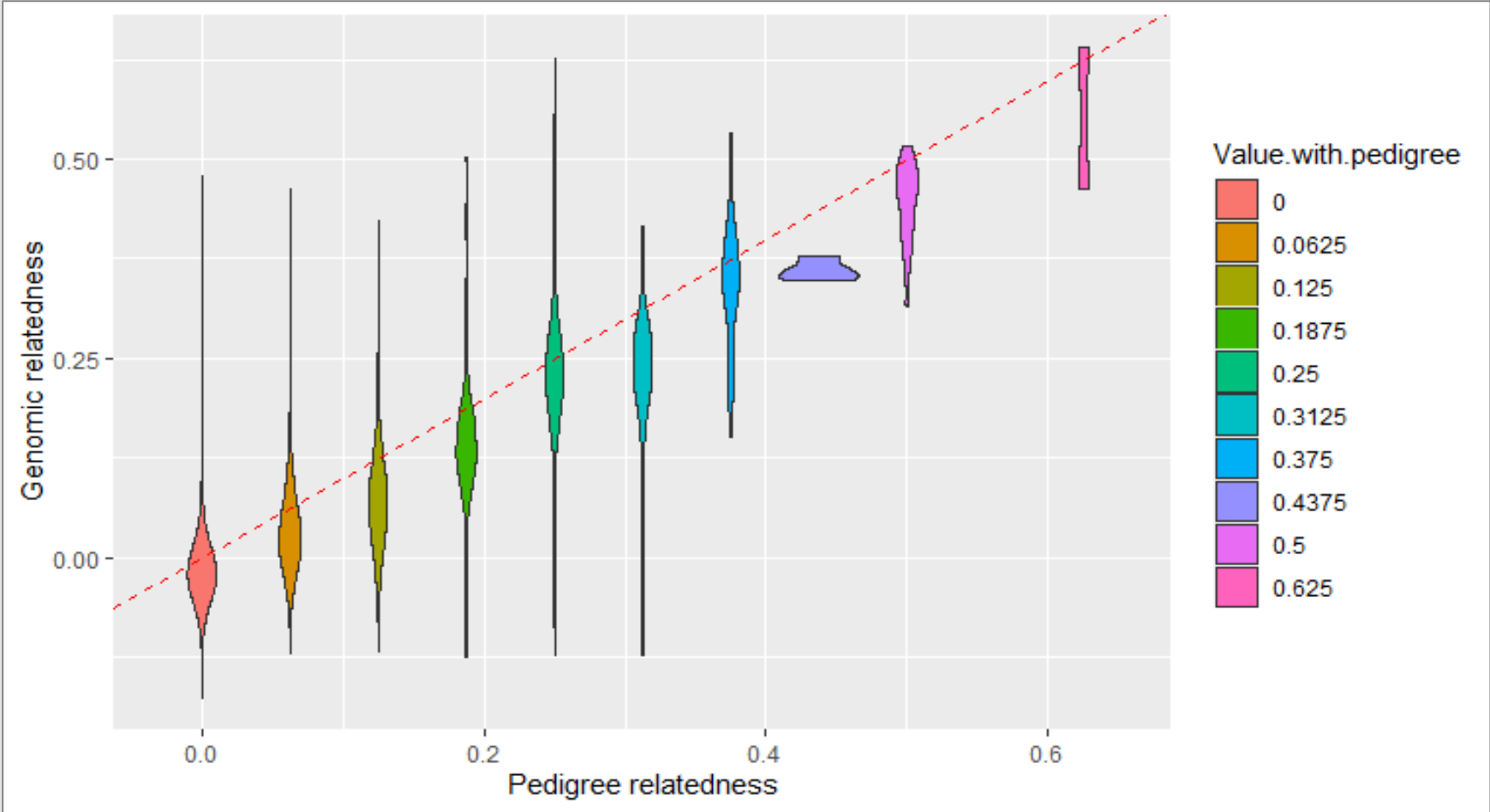
**A matrix**  
*calculated with pedigree information*



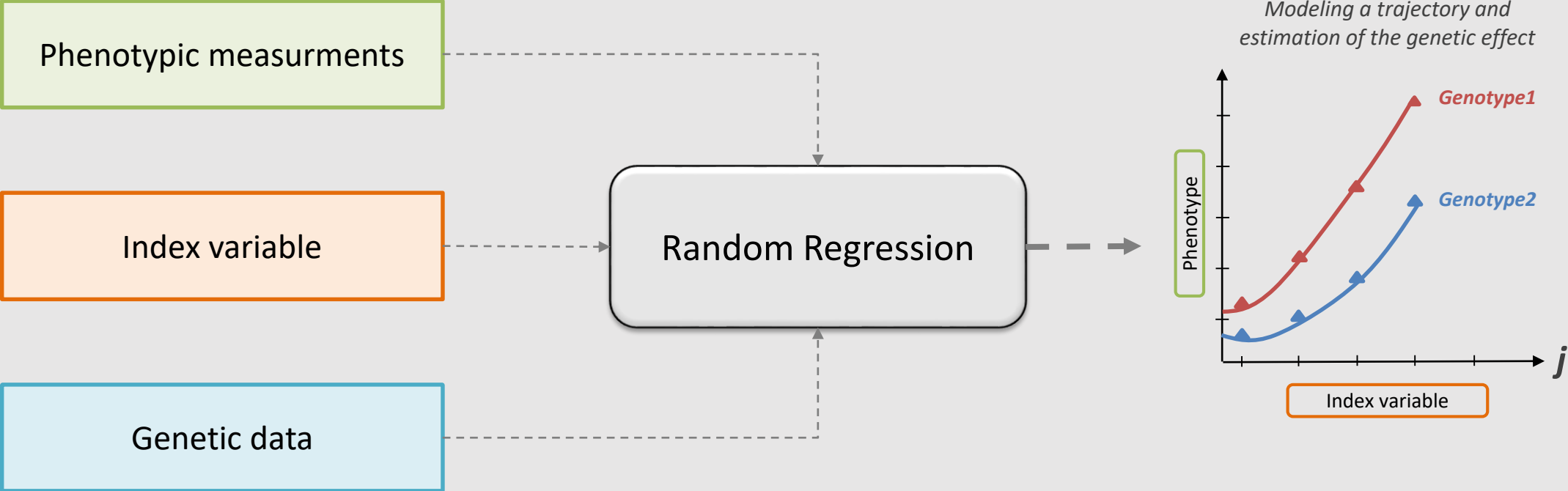
**G matrix (VR1)**  
*Calculated with 3000 SNPs information*



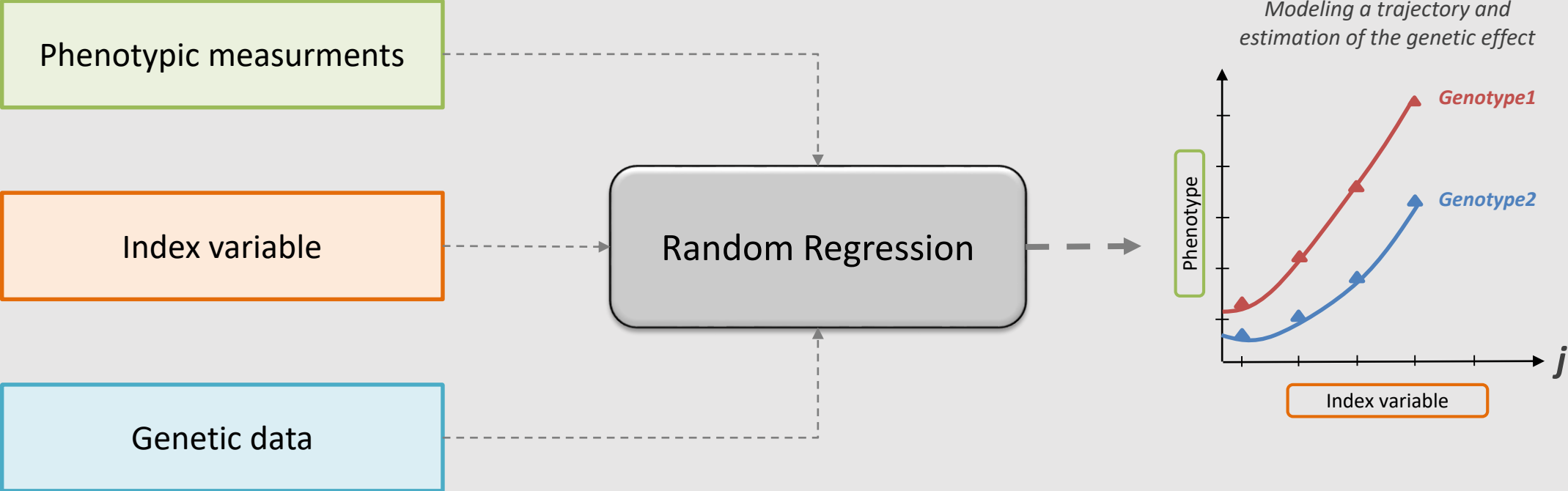
Comparison of relatedness calculated with genomic data and relatedness calculated with pedigree data



# Construction of norms of reaction

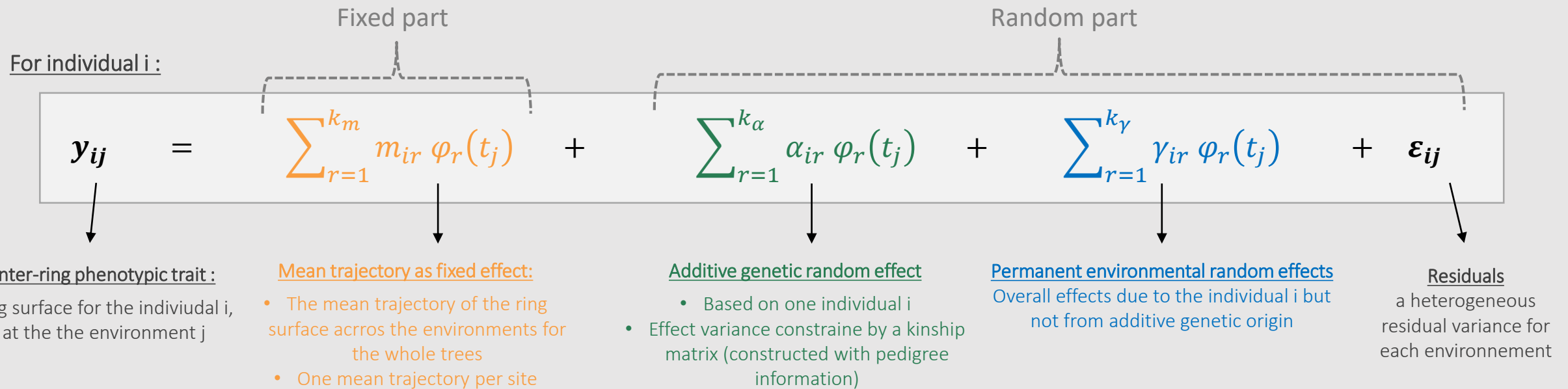
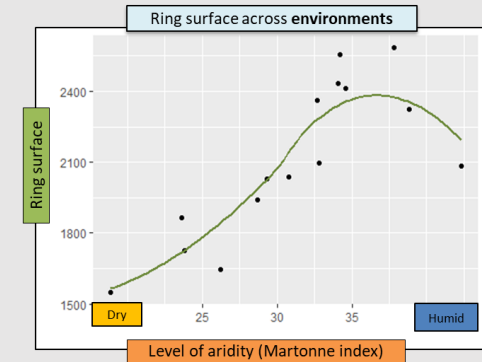


# Construction of norms of reaction



# Random regression model (RRM)

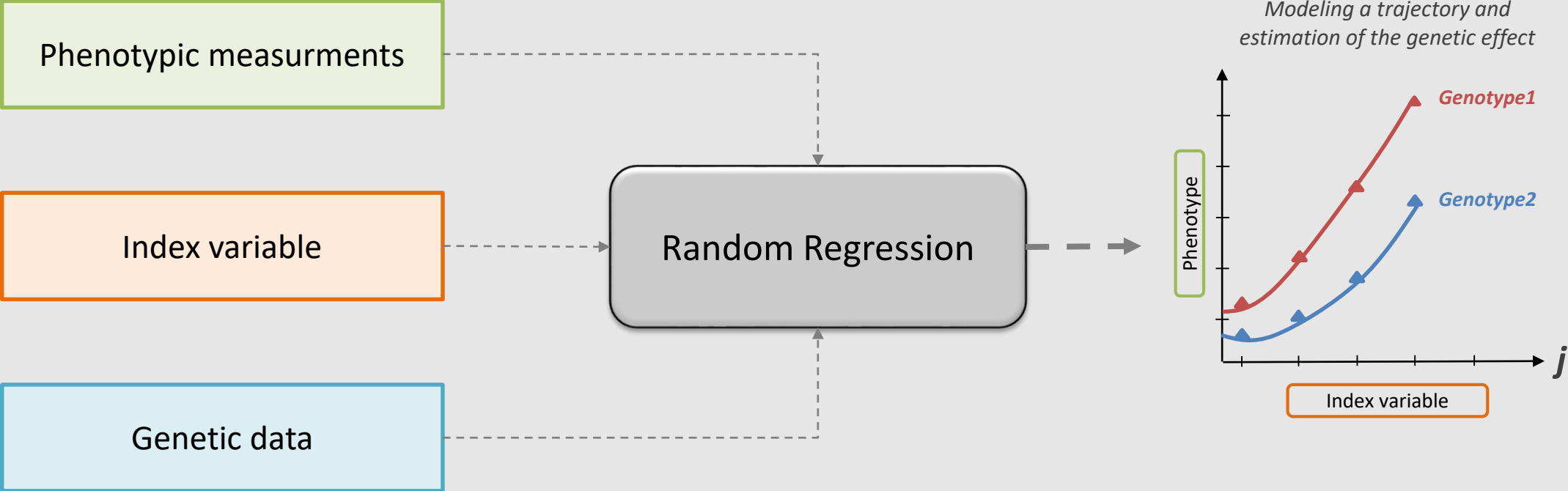
- We model individual trajectories according to the level of aridity of the environment (Martonne index)



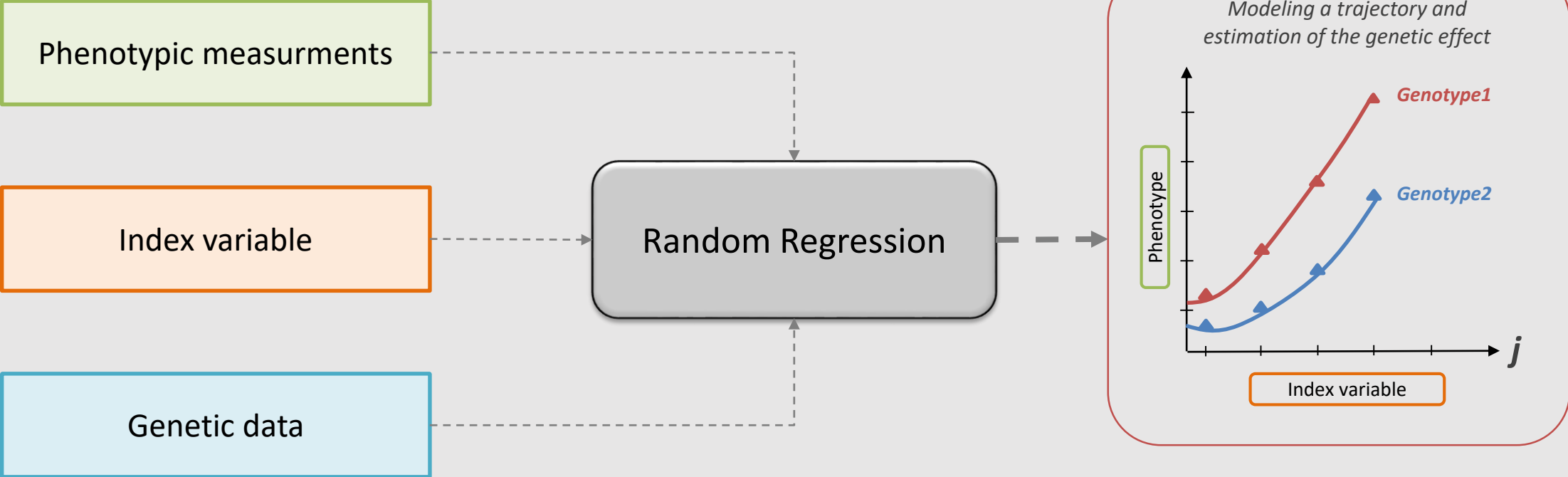
- Mean trajectories, additive genetic effect and permanent environmental effects are modeled by 2-order Legendre polynomials



# Construction of norms of reaction



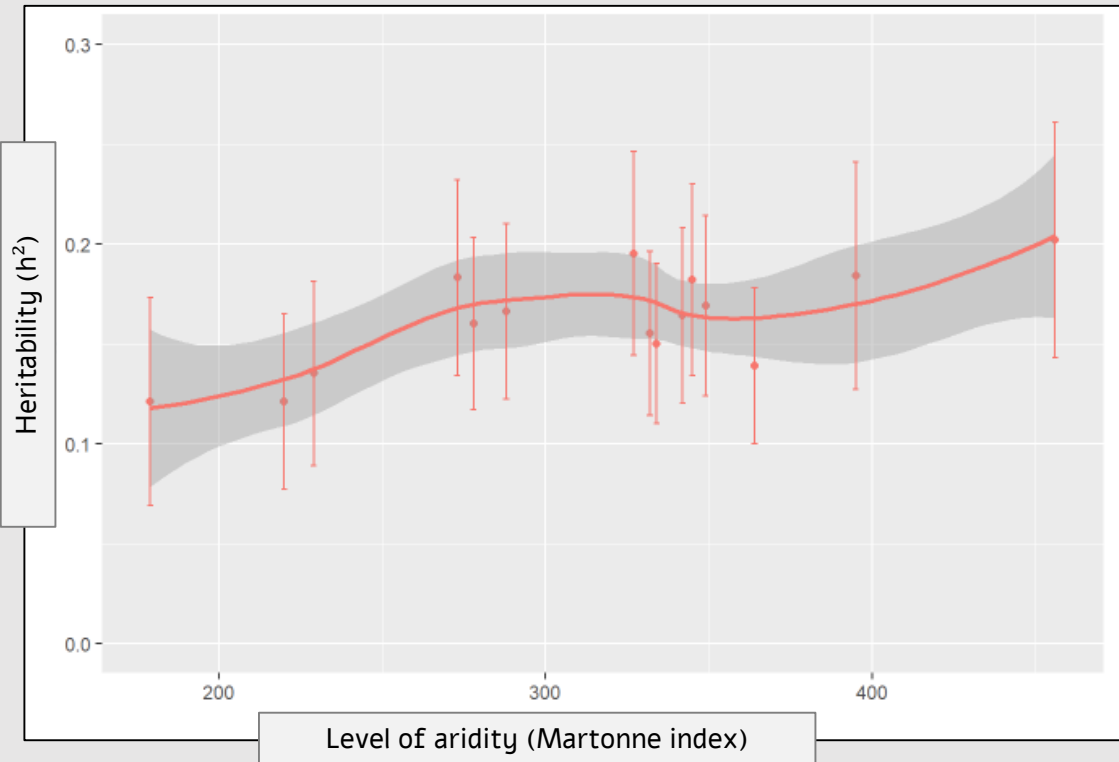
# Construction of norms of reaction



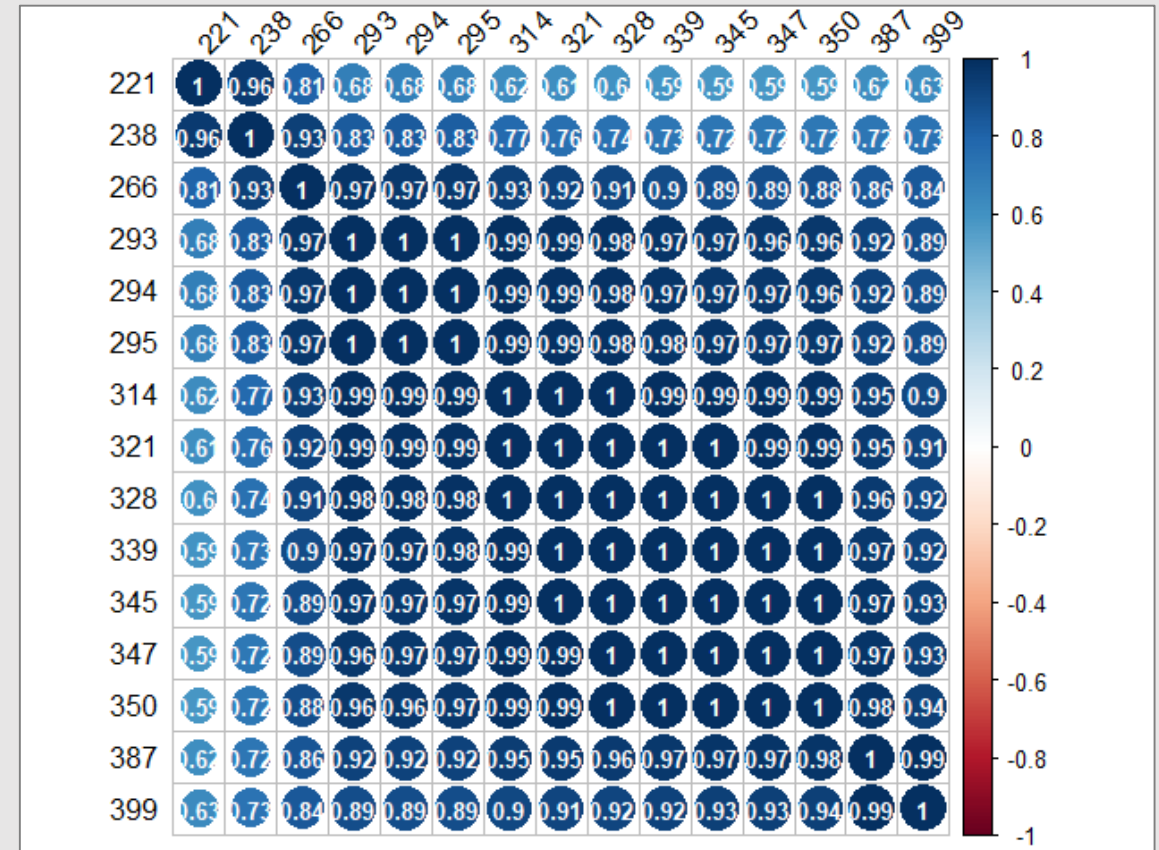
# Norms of reaction : heritability and genetic correlations

$$y_{ij} = \text{Mean trajectory } (t_j) + \text{Additive genetic effect}_i(t_j) + \text{Permanent environmental effect}_i(t_j) + \varepsilon_{ij}$$

Heritability of ring surface estimated with a random regression model of order 2 (RRM2)



Genetic correlations between environments (levels of aridity) estimated with RRM2



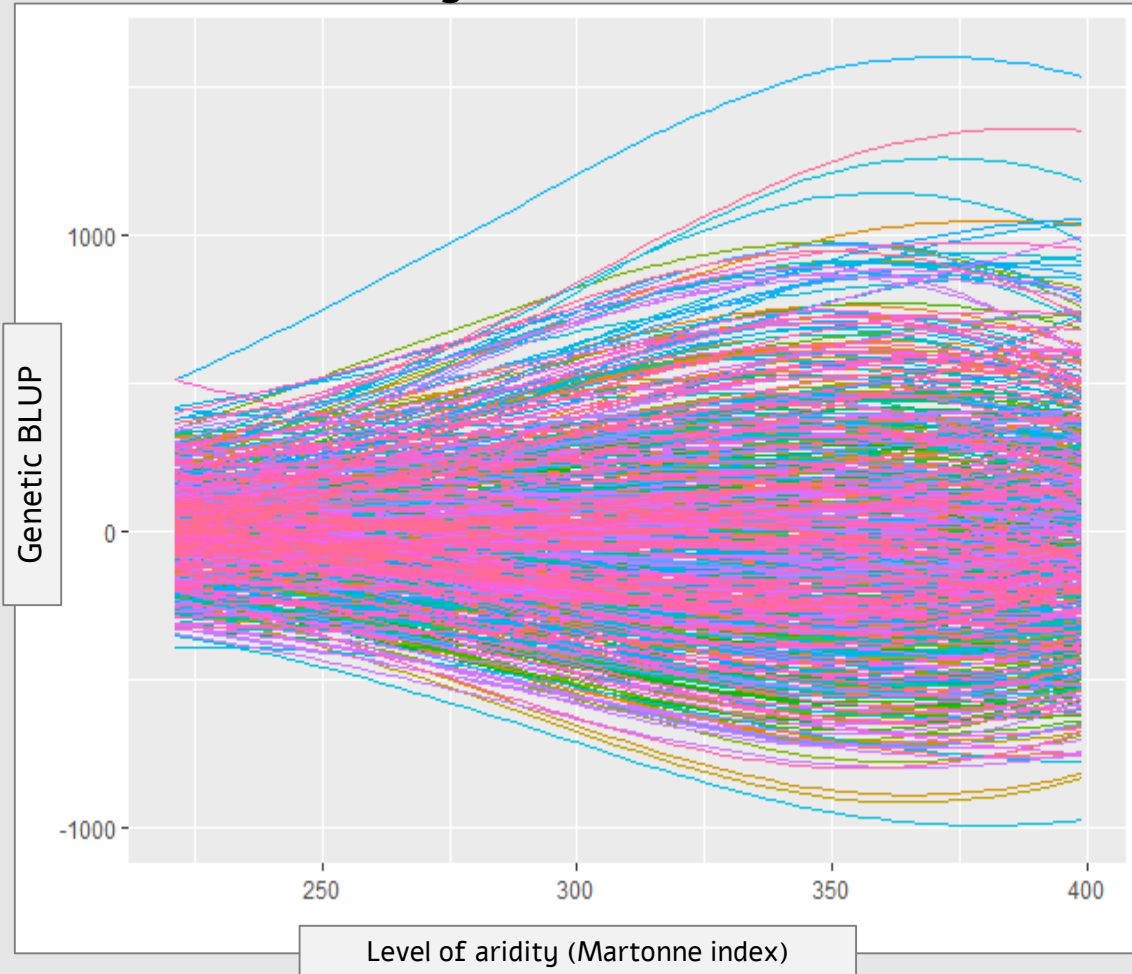
# Norms of reaction : additive genetic trajectories

$$y_{ij} = \text{Mean trajectory } (t_j) + \text{Additive genetic effect}_i(t_j) + \text{Permanent environmental effect}_i(t_j) + \varepsilon_{ij}$$

# Norms of reaction : additive genetic trajectories

$$y_{ij} = \text{Mean trajectory } (t_j) + \text{Additive genetic effect}_i(t_j) + \text{Permanent environmental effect}_i(t_j) + \varepsilon_{ij}$$

Evolution of individual genetic values across environments

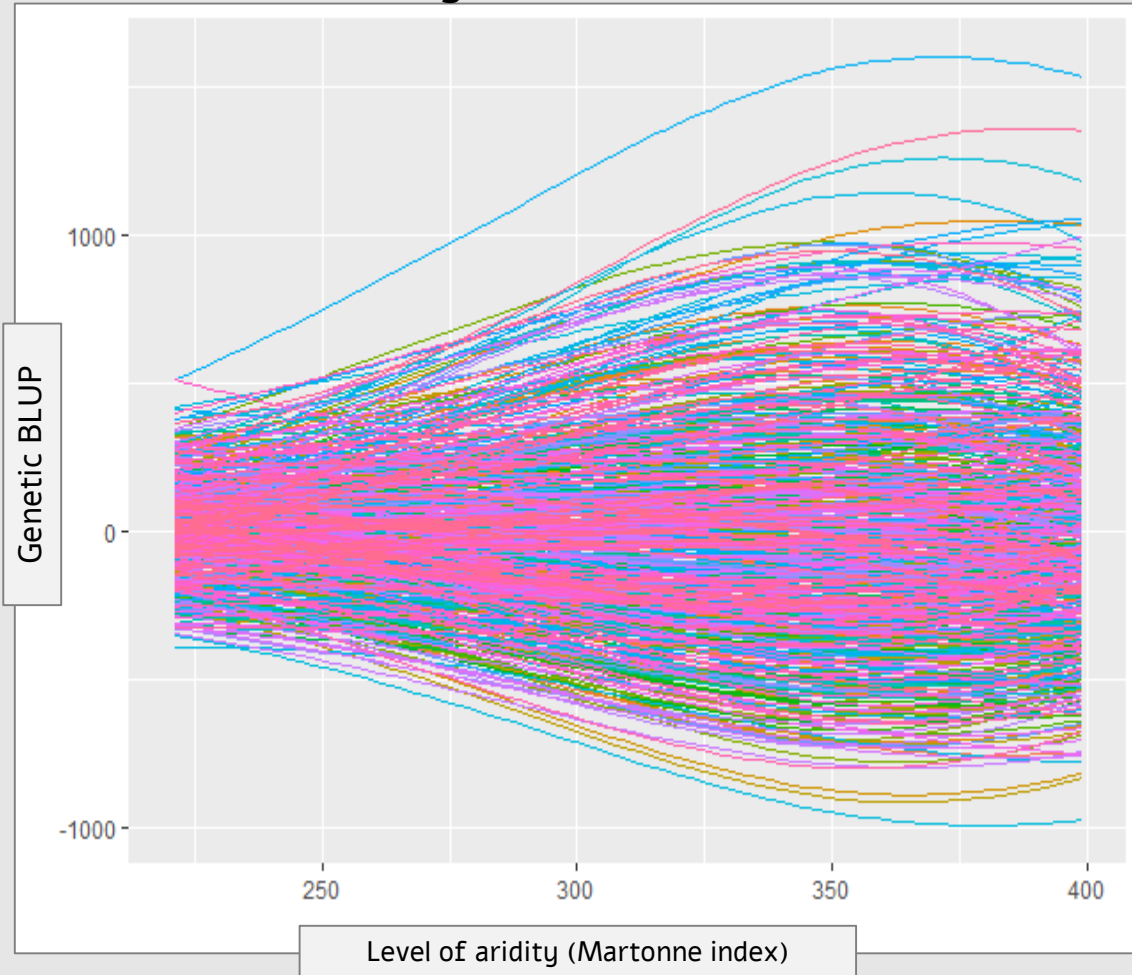


# Norms of reaction : additive genetic trajectories

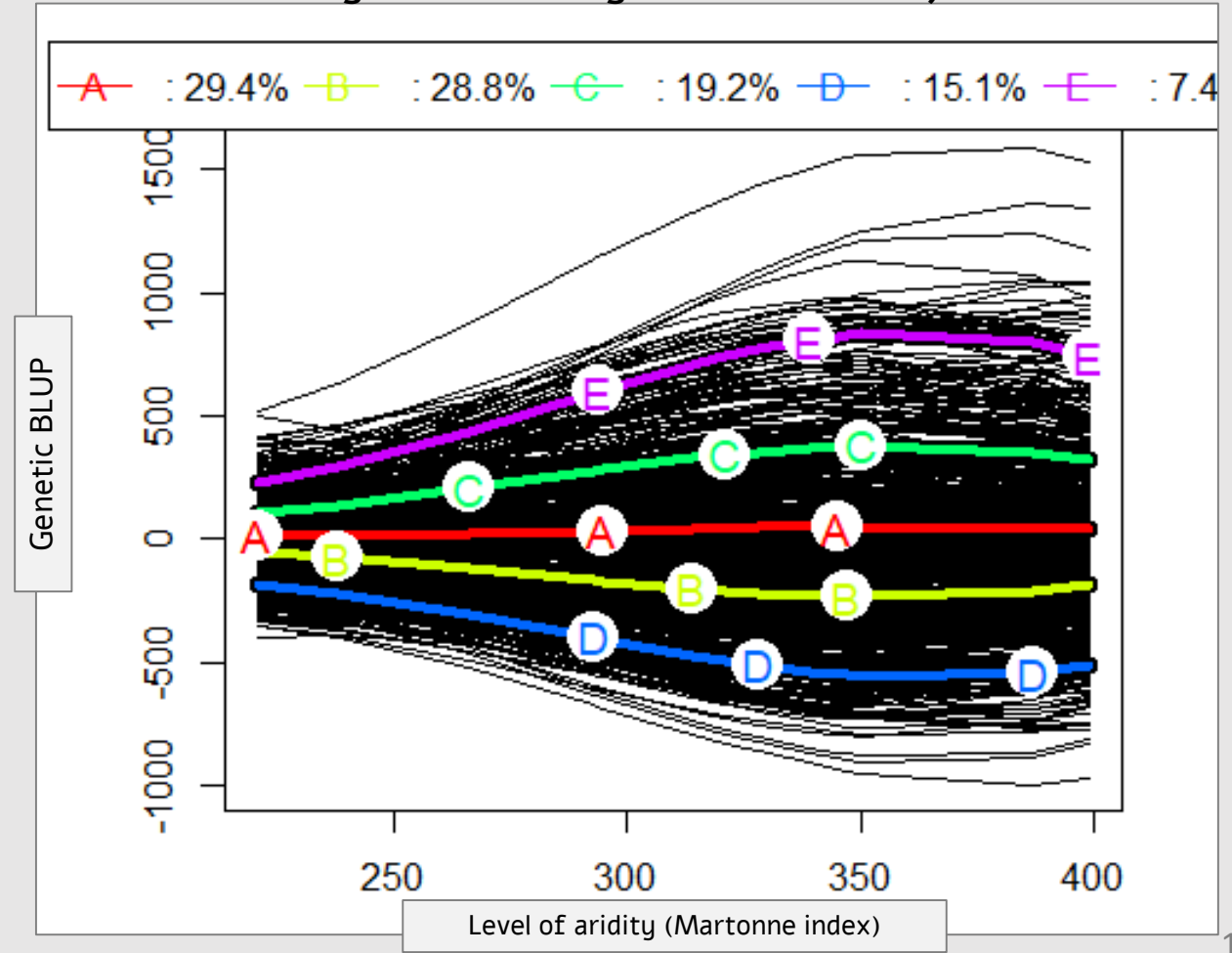
$$y_{ij} = \text{Mean trajectory } (t_j) + \text{Additive genetic effect}_i(t_j) + \text{Permanent environmental effect}_i(t_j) + \varepsilon_{ij}$$

Clustering of trajectories (Genolini et al. 2015)

Evolution of individual genetic values across environments



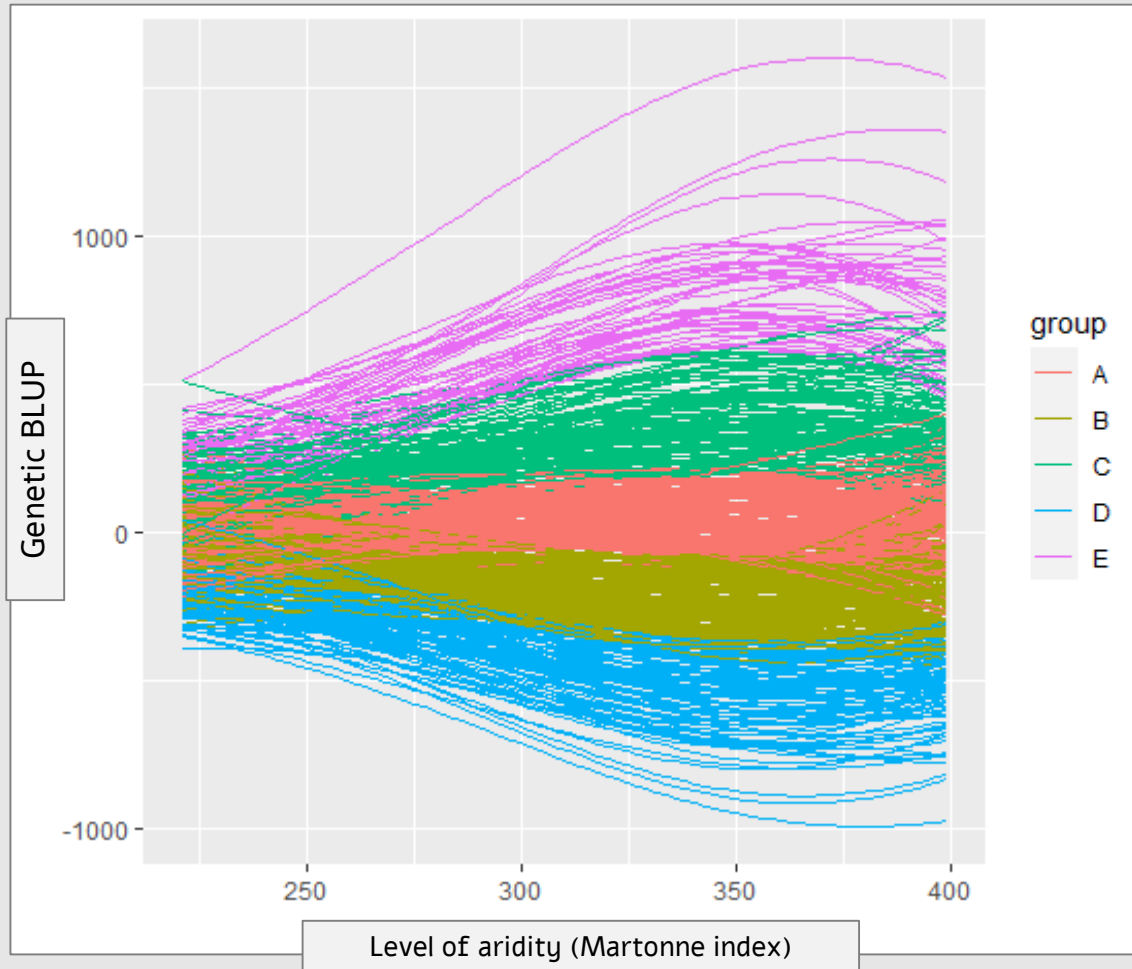
Clustering of individual genetic values trajectories



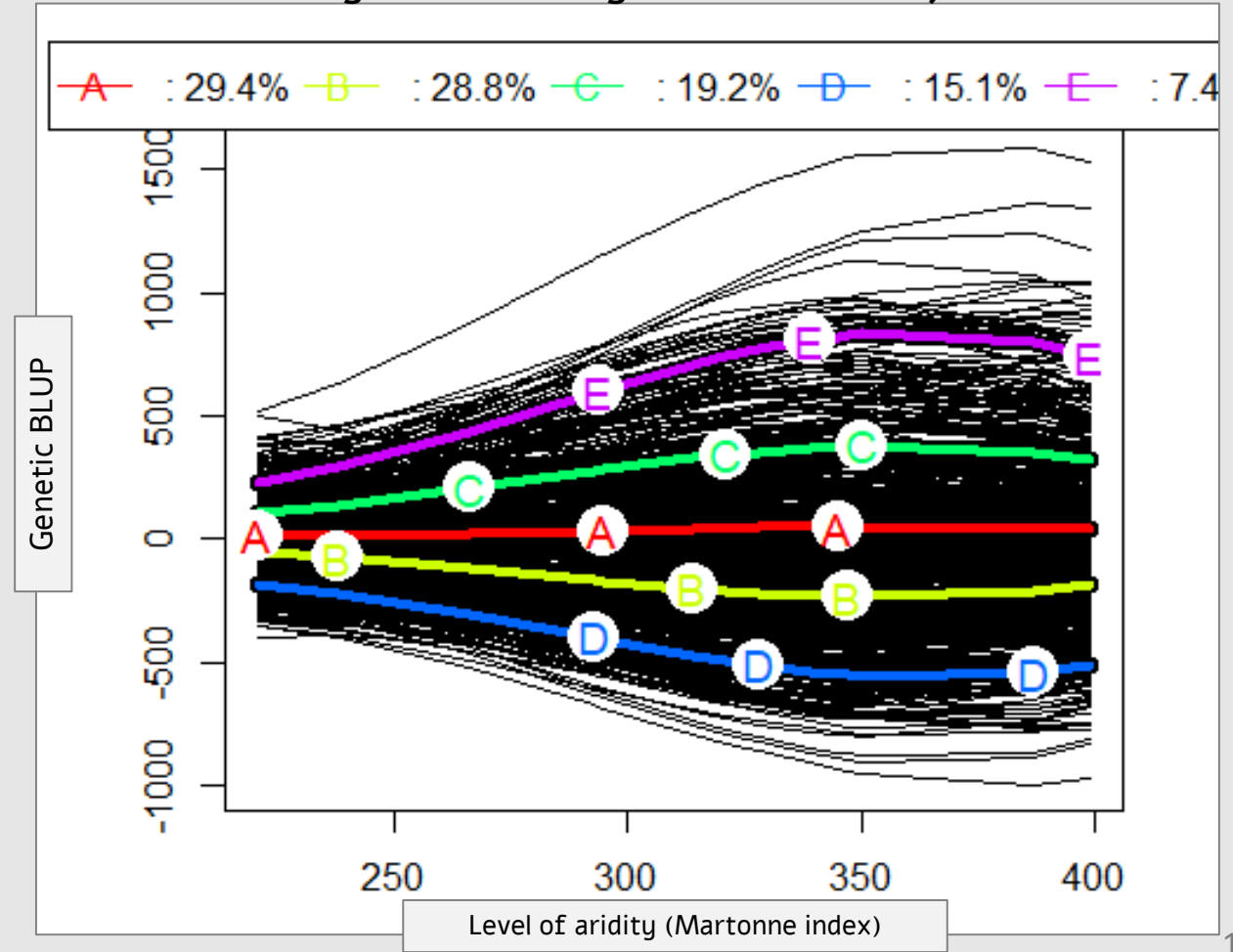
# Norms of reaction : additive genetic trajectories

$$y_{ij} = \text{Mean trajectory } (t_j) + \text{Additive genetic effect}_i(t_j) + \text{Permanent environmental effect}_i(t_j) + \varepsilon_{ij}$$

Evolution of individual genetic values across environments



Clustering of individual genetic values trajectories



# Conclusion and prospects

- With :

- Phenotypic measurements : densitometric profiles
- Simple climatic characterization : Martonne's aridity index
- Genetic data : pedigree or molecular markers information

→ Norms of reaction : individual genetic values across a range of environments

- Main prospects :

- Improve **environmental characterization** :
  - Consideration of previous years, better differentiation of experimental sites...
  - Use of a **growth simulation model** (GO+ : based on environmental characteristics of each site (stand density, groundwater height)) to incorporate more explanatory parameters into the annual index (soil water reserve, evapotranspiration demand)
- Extend to other **inter-ring phenotypic traits** (ex: average ring density)
- Construct NoR for **intra-ring growth**





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**Thank you for your attention**



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