

# **Reciprocal Recurrent Selection**

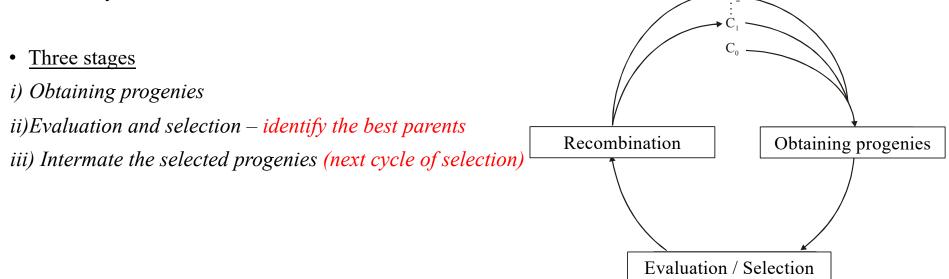
**Prof. Roberto Fritsche-Neto** 

roberto.neto@usp.br

Piracicaba, November 27<sup>th</sup>, 2019

### **Definition and scheme**

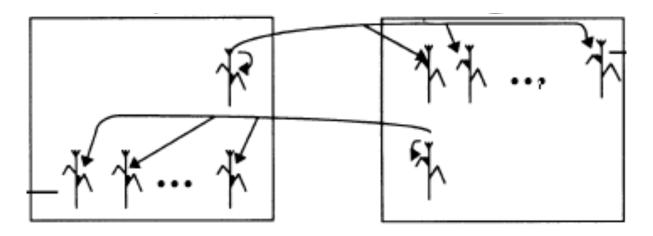
- Continuous process which aims the increasing of the allele frequencies but without miss substantial genetic variability.
- Dynamic process every cycle is possible to release na improved material and add more genetic variability



#### **General scheme of RRS** A x B and B x A (HS, FS or testers) $\mathbf{A} \mathbf{x} \mathbf{A} (\mathbf{S}_1 \text{ or } \mathbf{HS})$ $\mathbf{B} \mathbf{x} \mathbf{B} (\mathbf{S}_1 \text{ or } \mathbf{HS})$ $\mathbf{1}$ C<sub>o</sub> - Pop. A C<sub>o</sub> - Pop. B Х Intermate Intermate **Interpopulation hybrids** Pop. A Pop. B A x B and B x A Identify Identify **Evaluate / Select** the A parents of the B parents of the best hybrids the best hybrids Select hybrids **C**<sub>1</sub> - **Pop. A C**<sub>1</sub> - **Pop. B** $H = \left(p - r\right)^2 d$ Increase the heterosis between populations

## **Stages of RRS**

• Stage 1: obtaining progenies



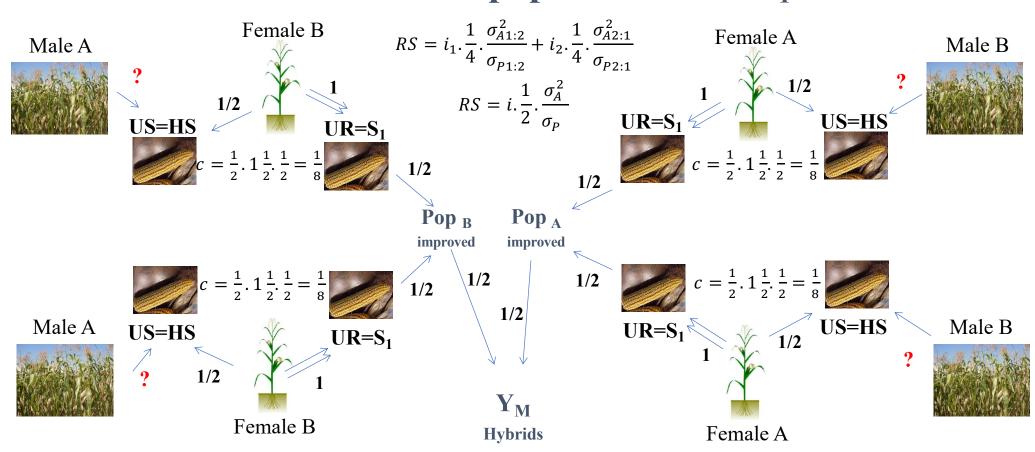
Testers, HS, or FS – Which one is the best to evaluate? HS or  $S_1$  – Which one is the best to intermate?

## **Stages of RRS**

- Stage 2: evaluation and selection
- Breeding objectives

i	Avaliação	Recombinação	c	Ne	Ne (10% de 200)	0/ 20 10 1 01
$RS = \frac{\iota}{\sigma_P} c \sigma_A^2$	MI	MI	1⁄4	4	80	$\begin{array}{cccccccccccccccccccccccccccccccccccc$
0 p	MI	$\mathbf{S}_1$	1⁄2	1	20	
$RS = i_1 \cdot c \cdot \frac{\sigma_{A1:2}^2}{\sigma_{P1:2}} + i_2 \cdot c \cdot \frac{\sigma_{A2:1}^2}{\sigma_{P2:1}}$	IC	IC	1⁄2	2	40	$N_{a} = 1$
	IC	$\mathbf{S}_1$	1⁄2	1	20	$Ive = \frac{1}{2F}$

- c = Parental control and additive covariance between the units of selection and recombination
- Effective population size evaluation (200) and intermate (10 to 20)
- Avoid to miss the genetic variability and boost the genetic drift



### **Scheme - interpopulation HS / S<sub>1</sub>**

# **Stages of recurrent selection**

- Stage 3: intermate each group separately
- Produce genetic variability for the next cycle
- Combine the superior allele/genes selected from different individuals in the newest genotypes

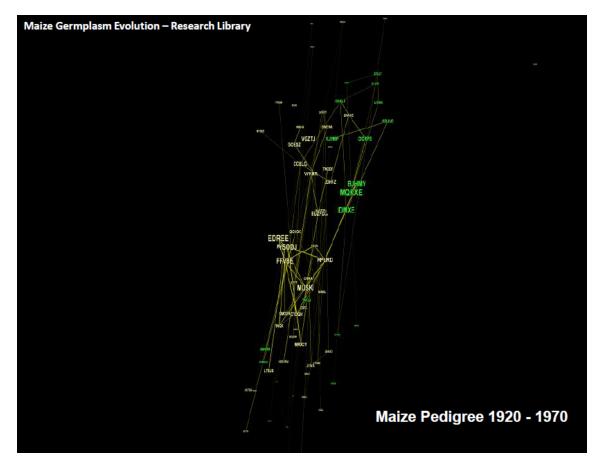
### **Ireland Method**



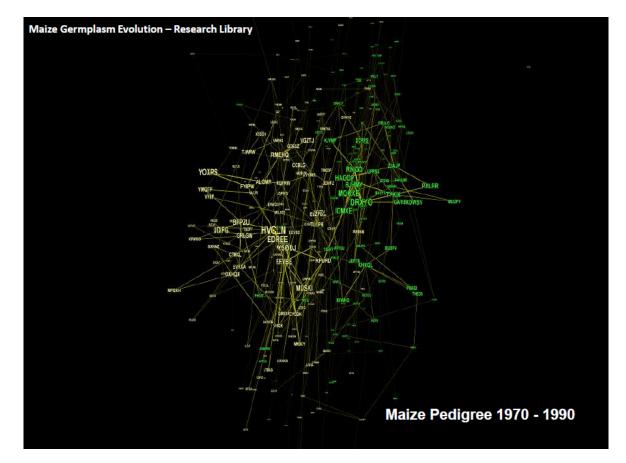
- 50 plants at least in the female rows
- Just one cycle of random intermate is enough to achieve the HWE
- Use the same number of seeds to hybridize and to compose the post-harvest sample

2N = number of gametes used  $u_0 =$  mean of gametes per parent  $\sigma =$  variance for number of gametes

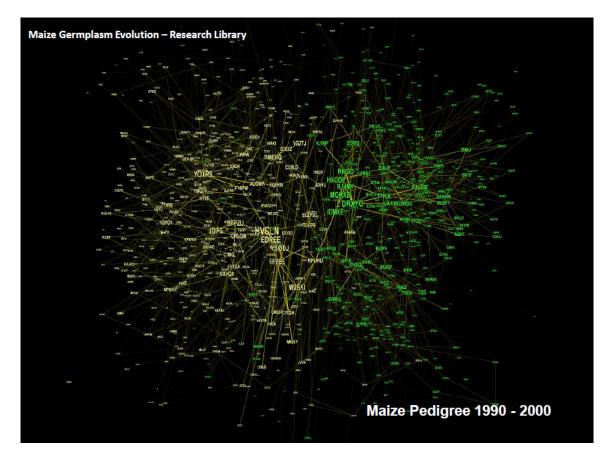
# **Germplasm of DuPont - Pioneer**



# **Germplasm of DuPont - Pioneer**

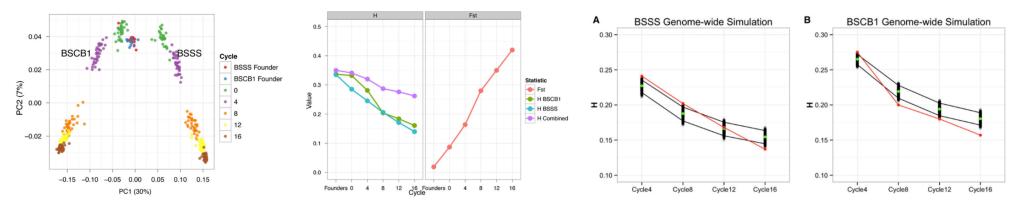


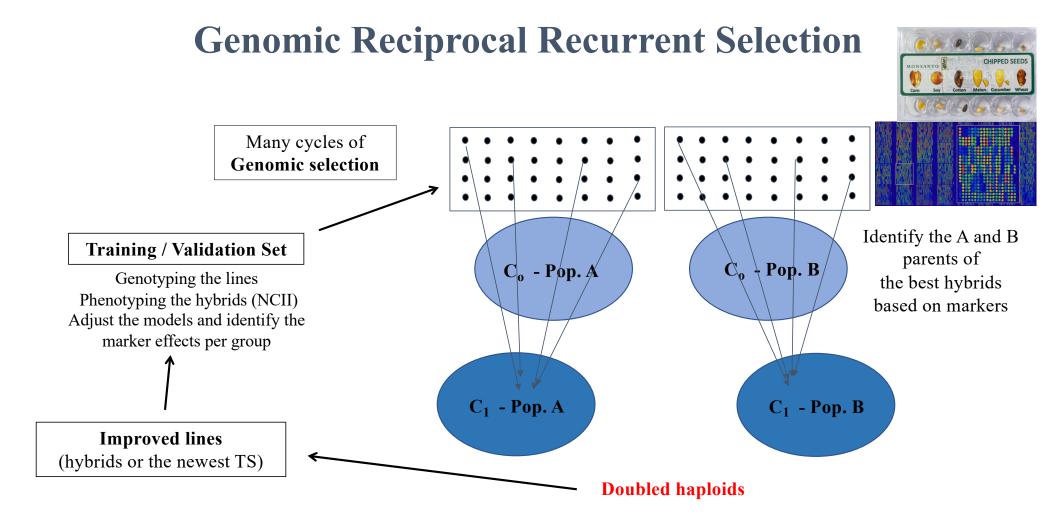
# **Germplasm of DuPont - Pioneer**



### **Drift and selection in RSS**

- Gerke et al (2015)
- Iowa RRS program started in 1949 16 cycles
- Founders: BSSS population 16 and BSCB1 12 inbred lines
- 100 progenies HS/S1 (Ne = 1 each one)
- Between cycle 4 and 8, 2 self generations
- Using empirical and simulated data they evaluated Fst, Ht, diversity, and distance





# Long-term selection gains in GS

 $C = \frac{1}{2} \left( 1 + \frac{1}{n_m} X X^T \right)$ 

RS = xTa

- Gorjanc et al (2018)
- Coanscestry = C = IBD
- where, X = M 1 (0, 1, 2) c = xTCx  $x = \frac{1}{2nc}n$
- Expected inbreeding
- where, n vector of number contributions (0, 1,2 3, or 4)
- and n<sub>c</sub> number of crosses

