



# Monitoring small poultry populations using simulated annealing algorithms

Hervé Chapuis, Daniel Guemene, Nabeel Alnahhas

## ► To cite this version:

Hervé Chapuis, Daniel Guemene, Nabeel Alnahhas. Monitoring small poultry populations using simulated annealing algorithms. 10. European Symposium on Poultry Genetics (ESPG), Jun 2017, St Malo, France. 2017. hal-02786281

**HAL Id: hal-02786281**

**<https://hal.inrae.fr/hal-02786281>**

Submitted on 4 Jun 2020

**HAL** is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers.

L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.



Daniel GUÉMENÉ<sup>1</sup>, Nabeel ALNAHHAS<sup>1</sup>, Hervé CHAPUIS<sup>2</sup>

<sup>1</sup> SYSAAF, Centre INRA Val de Loire, URA, 37380 Nouzilly, FRANCE

<sup>2</sup> UMR 1388 INRA-INPT GenPhySE, 31326 Castanet-Tolosan Cedex, FRANCE



GenPhySE



## Introduction

Breeding schemes have to produce immediate genetic gain without impairing genetic variability, in order to preserve future gains.

Two strategic steps require attention:

- **Breeder selection.** Results from an equilibrium between average EBV and coancestry of selected individuals.
- **Mating plan design.** Aims at achieving minimization of expected inbreeding in the next generation.

In poultry breeding, where schemes are built under a hierarchical mating design, genetic contributions are both equal and discrete : once selected, each sire is mated with  $d$  dams, and egg collection period length is identical for all females.

**These characteristics are stressed in the case of small populations, which may be local breeds or experimental populations.**

Simulated Annealing (SA) methods prove to be effective tools in order to achieve :

- Maximization of genetic level of selected breeders, subject to a constraint on their average coancestry.
- Minimization of average coancestry of selected breeders, subject to a constraint on their average EBV.
- Minimization of expected coancestry in the next generation, allowing or not sisters to be mated to the same male.

## Material & Methods

### Simulated Population

Demographic parameters: 15 [(1♂ × 3♀) → 3 × (3♂ + 3♀)]

2 traits :

1. MAIN OBJECTIVE (e.g. growth)  $h^2 = 0,4$  ← TO BE MAXIMIZED

$$\rho_G = \rho_E = -0,25$$

2. ANCILLARY OBJECTIVE (e.g. egg number)  $h^2 = 0,2$  ← TO BE CONSTRAINED TO ZERO

3 methods for breeder selection:

1. **REF** : 1 male offspring /sire & 1 female offspring /dam are selected among candidates exhibiting an EBV > -0,5  $\sigma_G$  for the ANCILLARY TRAIT.
2. **BLUP50** : maximization of the MAIN OBJECTIVE, while keeping the ANCILLARY TRAIT equal to zero and aiming at and inbreeding trend equal to half the value obtained in the case of an unconstrained selection.
3. **INBREDMINT** : minimization of coancestry for selected animals, while aiming at achieving: 1) a gain on the MAIN OBJECTIVE equal to what is obtained with REF, and 2) a null trend for the ANCILLARY TRAIT.

3 mating strategies:

1. Random.
2. Optimized and unconstrained → minimization of expected inbreeding.
3. Minimization of expected inbreeding BUT two related dams (full-sibs sisters) can't be mated to the same sire.

### Indicators

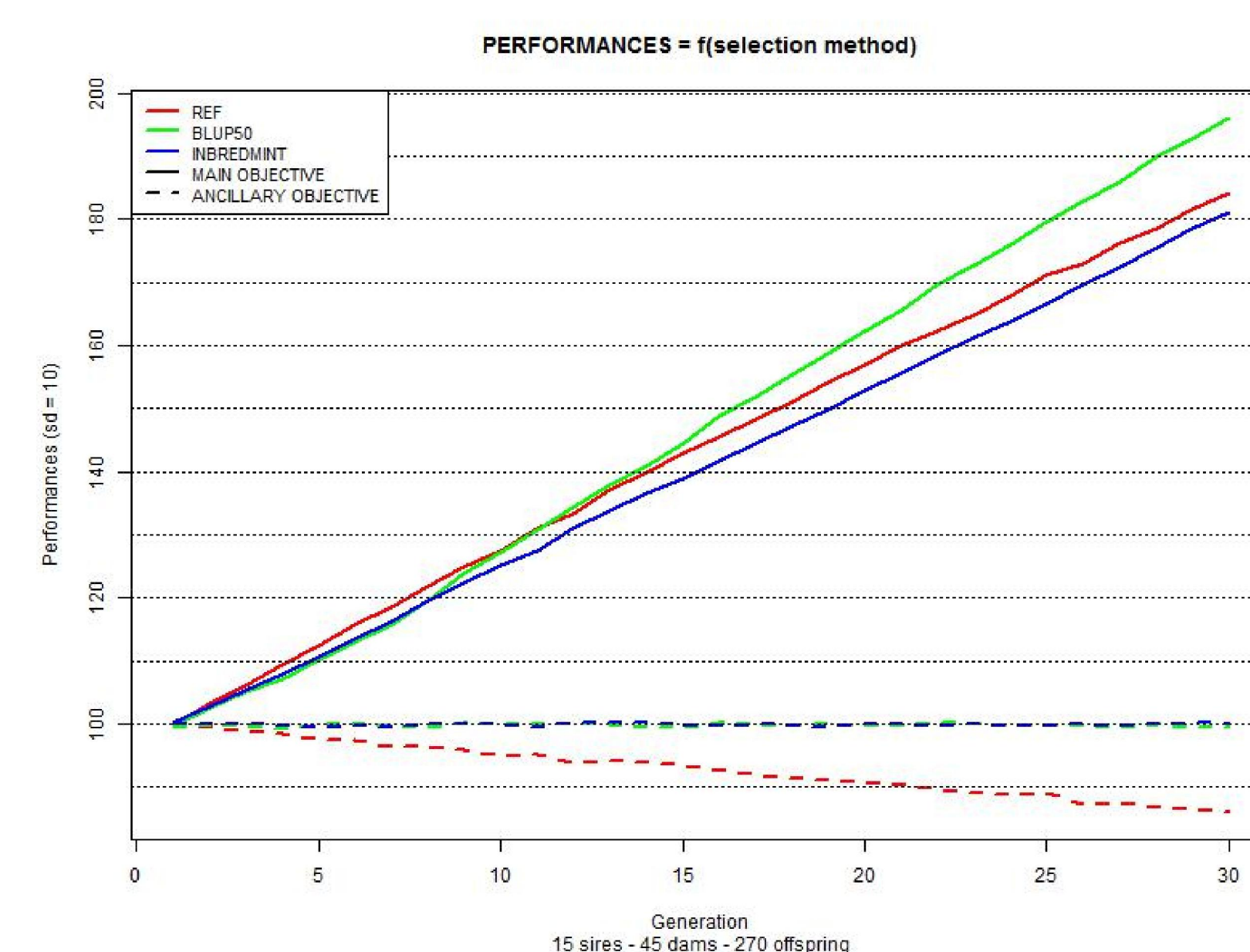
Average EBV, average coancestry, effective populationsize ( $N_e$ ).

(« démographic  $N_e$  » = 45)

## Conclusions

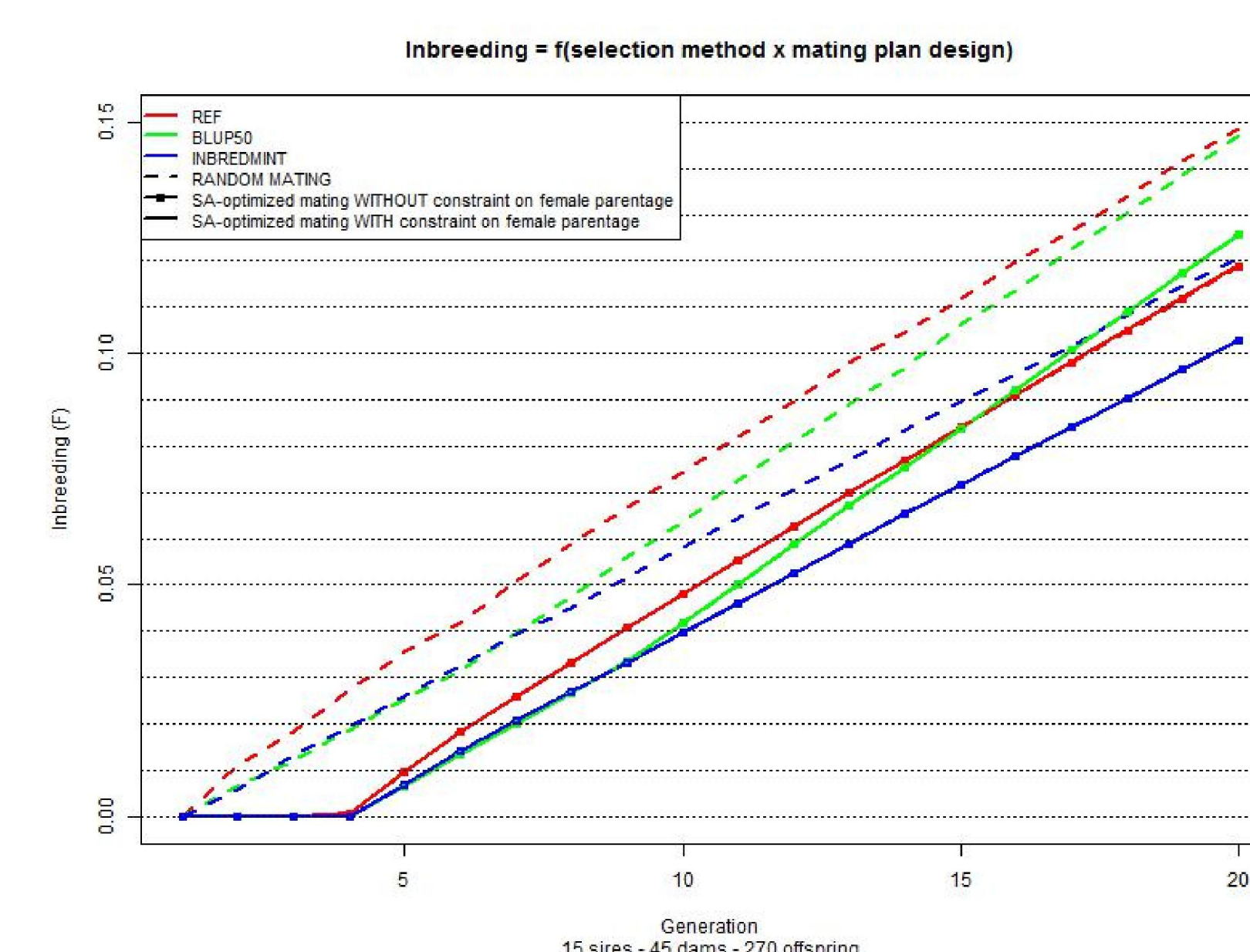
- SA methods proved to be effective tools in order to monitor small populations.
- Further studies should be carried out in the case of selection without individual cages, when pedigree is only known *a posteriori*, after parentage assignment.
- An additional optimization will certainly be required, in order to insure the preservation of the marker panel assignment power.

## Results



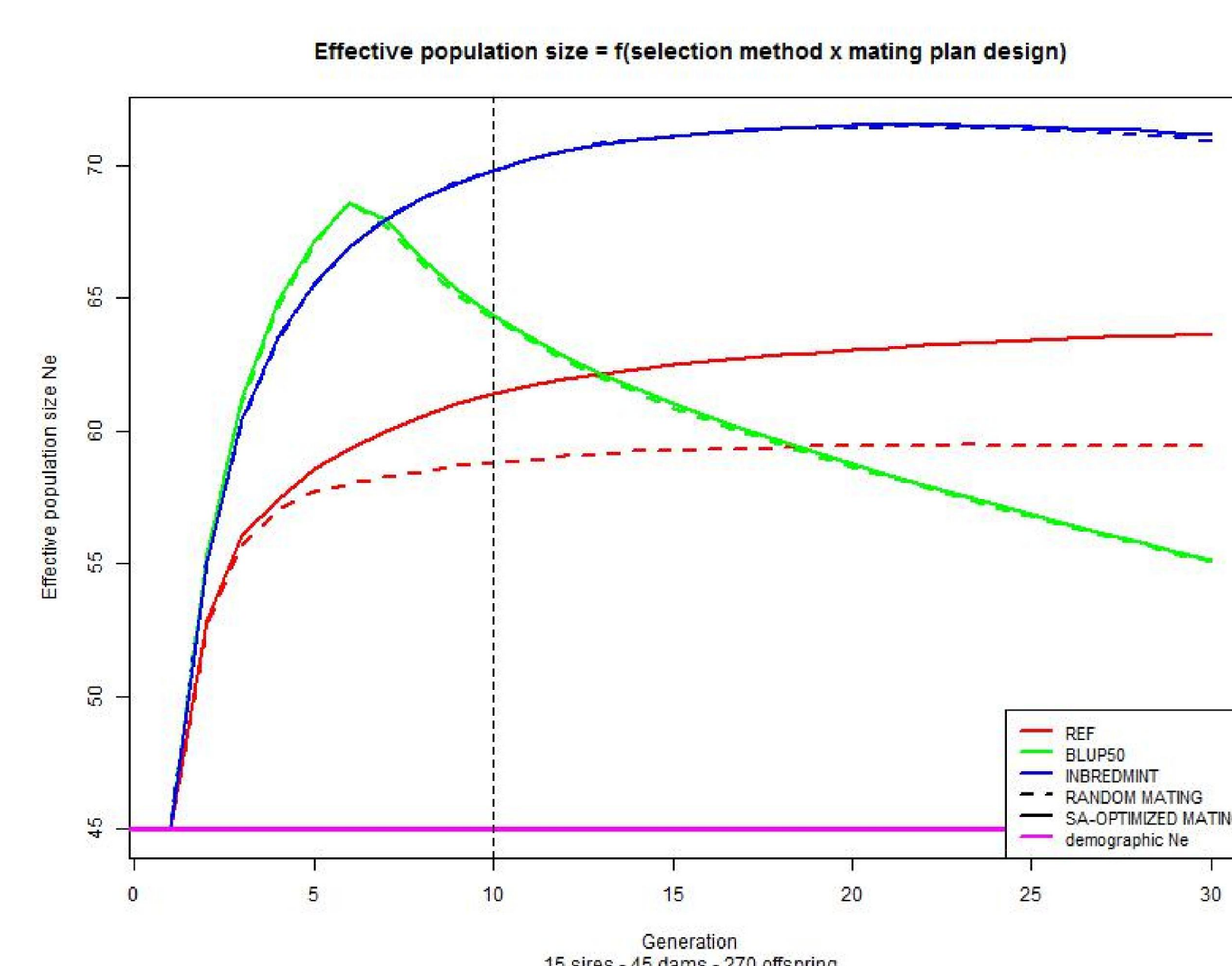
### SELECTION RESPONSE

- Only SA methods can achieve the desired null trend for the ANCILLARY TRAIT.
- In the short term, **REF** provides a better gain on the MAIN OBJECTIVE.
- Beyond 10 generation, **BLUP50** clearly outperforms **REF**.



### INBREEDING

- SA methods clearly outperform **REF**.
- Mating optimization does not affect the slope of the inbreeding trend.
- The main effect is on the intercept, as it delays the onset of inbreeding.
- Preventing the mating of related females to the same sire is ineffective (the two curves overlap).



### EFFECTIVE POPULATION SIZE

- **INBREDMINT** leads to the highest value of  $N_e$ . **REF** remains stable.  $N_e$  decreases with **BLUP50**.
- The mating design has no effect on  $N_e$  when SA methods are used for breeder selection.
- No reliable conclusion should be drawn from pedigree data when less than 7-10 generations are available .