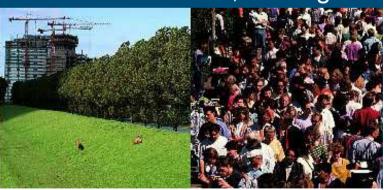
The current situation of regional cattle breeds within the Netherlands

Myrthe Maurice-Van Eijndhoven

E-mail: myrthe.maurice-vaneijndhoven@wur.nl

Centre for Genetic Resources, the Netherlands Animal Breeding & Genomics Centre Abstract number 3243 Session 8 EAAP 2008, 24 August







Background

- Cattle farming over the last decades
 - From dual purpose (beef and dairy) to more specialized farming
 - Use of highly selected bulls
 - Introgression of the Holstein Friesian (HF) cattle in the early seventies
- Consequences for local breeds
 - Demographic
 - Genetically



Research questions

1 Changes in population size in the MRY and the FH cattle breed populations over the last three decades?

Changes in genetic diversity in MRY and FH cattle breed populations over the last three decades?

Which part of the total genetic diversity of the FHbreed is already stored in the genebank?





Methods

Data

Pedigree data with 4,446,561 MRY- and 7,318,522 FH (purebred + crossbred)

- Estimations
 - Average inbreeding
 - Average coancestry
 - Optimal contributions

NB: unknown parents → bias in estimations

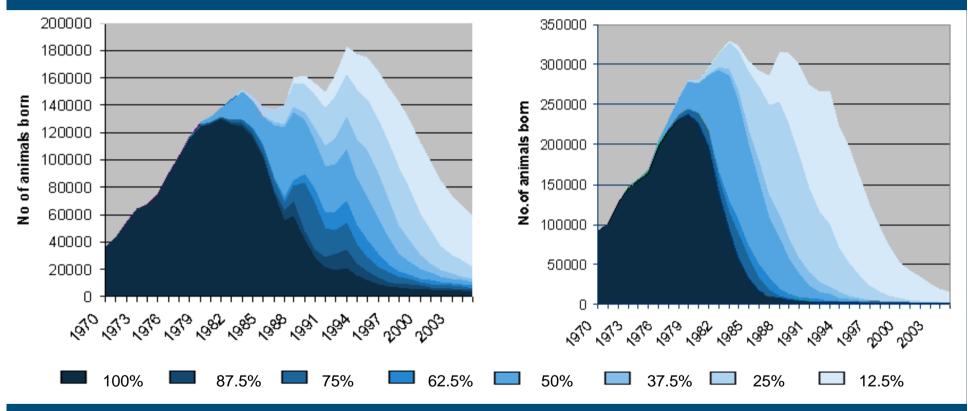


Results: Demographics (1)



Consequences on development population sizes 1

MRY FH



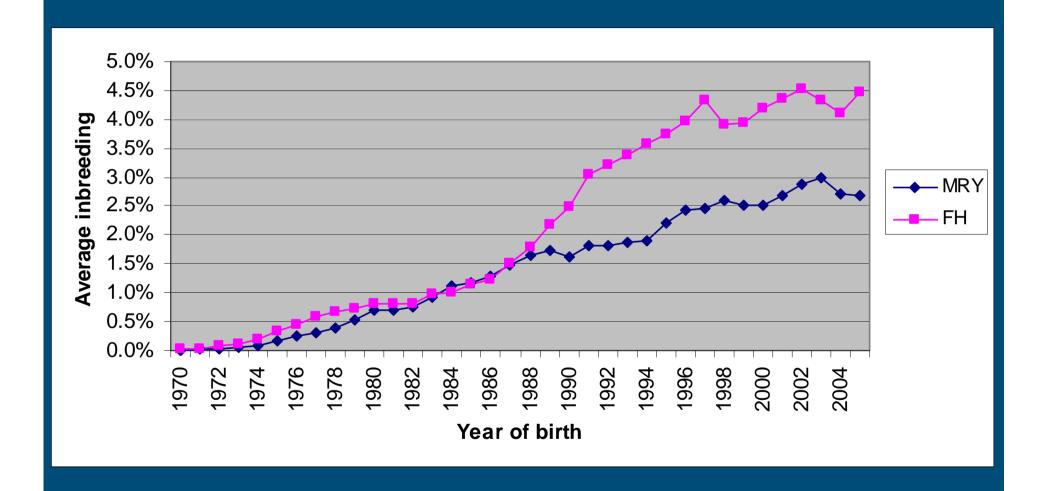
% own genes



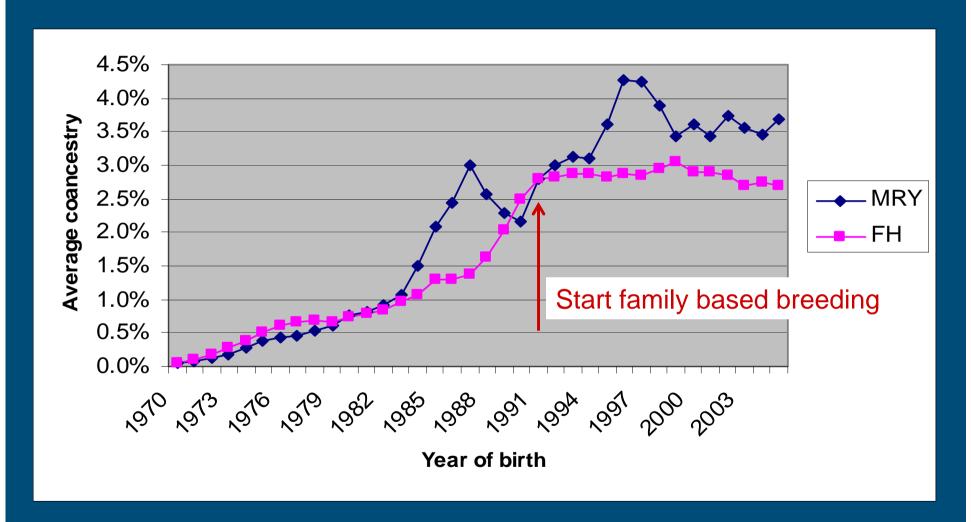
- The purebred population size strongly decreased for both breeds
 - > 99% in the last three decades
- Some arguments of farmers:
 - Milk production level
 - Breed characteristics: robustness; sober; recognizable; meat production and milk composition

Results: Genetic diversity (2)

Average inbreeding



Average coancestry



2

- Genetic diversity decreased in both populations;
- The 'family based' breeding structure of FH had a positive effect on the conservation of genetic diversity;
- Utilization of a few bulls causes a strong decrease in genetic diversity.

Results: Gene bank collection (3)



 Using the optimal contribution method a maximum of genetic diversity can be reached by selecting 91 bulls.

62 of these 91 selected males were bulls for which semen is stored within the gene bank. These 62 gene bank bulls cover 75% of the total calculated sire contributions.



Conclusion: FH gene bank collection

3

The CGN genebank collection is an important genetic resource of FH- material

Final messages

- Farmers and breeding organisations need to be aware of the effect of bull selection
- Preservation of breeds in a genebank alone cannot prevent breeds from extinction by itself
- Give attention to special breed characteristics



Thank you for your attention



