Optimizing design of small-sized nucleus dairy cattle breeding programs with minimal recording

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Introduction

 Current genetic improvement in developing countries is through semen importation



Temperate



Tropical

 GxE between regions estimated to be 0.49 (Ojango and Pollot, 2002)



Environment specific breeding programs

- Alternative approach: environment-specific breeding programs
- Limitation minimal and erratic pedigree and performance recording at farm level (Wasike et al., 2011)
- Negatively impacts genetic evaluation of selection candidates



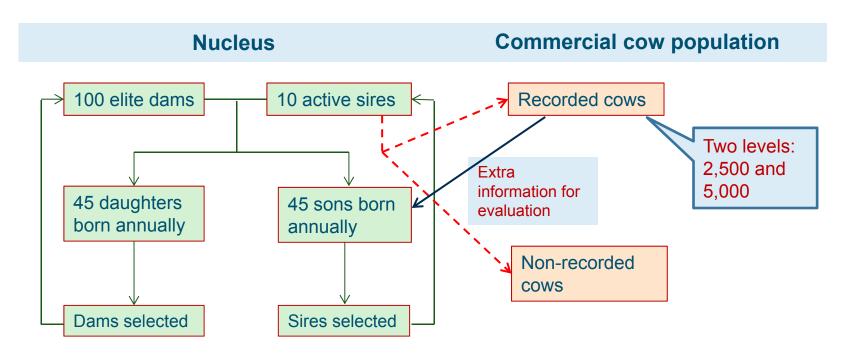
Objective

- Small-sized nucleus dairy cattle breeding program?
- What selection strategy to adopt?
 - Response to selection
 - Accuracy
- We ignored inbreeding for this study



Materials and Methods

Deterministically simulated a nucleus program



SelAction (Rutten et al., 2004)



Materials and Methods

Selection strategies

Abbreviation	Description	
DP	Phenotypes of nucleus dams	
PT	Progeny testing	
GS	Genomic selection	
GS+DP	Genomic + nucleus dams	
GS+PT	Genomic + progeny testing	



Materials and Methods

- GS was implemented by mimicking a correlated trait with $h^2 = 1$ and genetic and phenotypic correlations were calculated following Dekkers (2007)
- Selected for a single trait total merit trait
- Truncation selection with 8 age-classes



Results – response to selection (ΔR)

Response for the basic DP scheme in genetic standard deviation (σ_a)

	Number of CRC		
Scheme	2,500	5,000	L (years)
DP	0.042	0.047	4.2

 For comparison alternative selection strategies were benchmarked against basic DP scheme and presented as a percentage



Results – response to selection (ΔR)

 ΔR Number of CRC 2,500 L (years) Scheme 5,000 13.5 6.2 PT 27.0 GS 24.3 70.3 3.0 4.2 GS+DP 24.3 43.2 GS+PT 16.2 6.2 29.7

Extra response as percentage of the response in DP schemes

For these results the pedigree is assumed to be known accurately



Results – accuracy (r_I)

	r_I		
	Number of CRC		
Scheme	2,500	5,000	
PT	0.62	0.73	
GS	0.21	0.30	
GS+DP	0.33	0.39	
GS+PT	0.64	0.74	



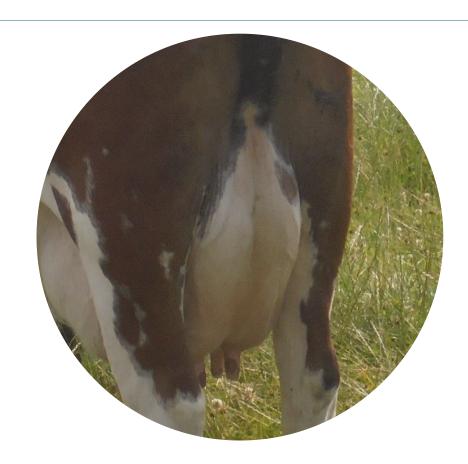
Conclusions

 Feasibility exists for creating genetic gains through nucleus programs with minimal performance recording

- GS will have the highest responses
- Lower accuracies in GS will be offset by the higher annual responses



Thank you for your attention!



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