

Potential benefits of distributing males and females among phenotyping candidates in genomic selection

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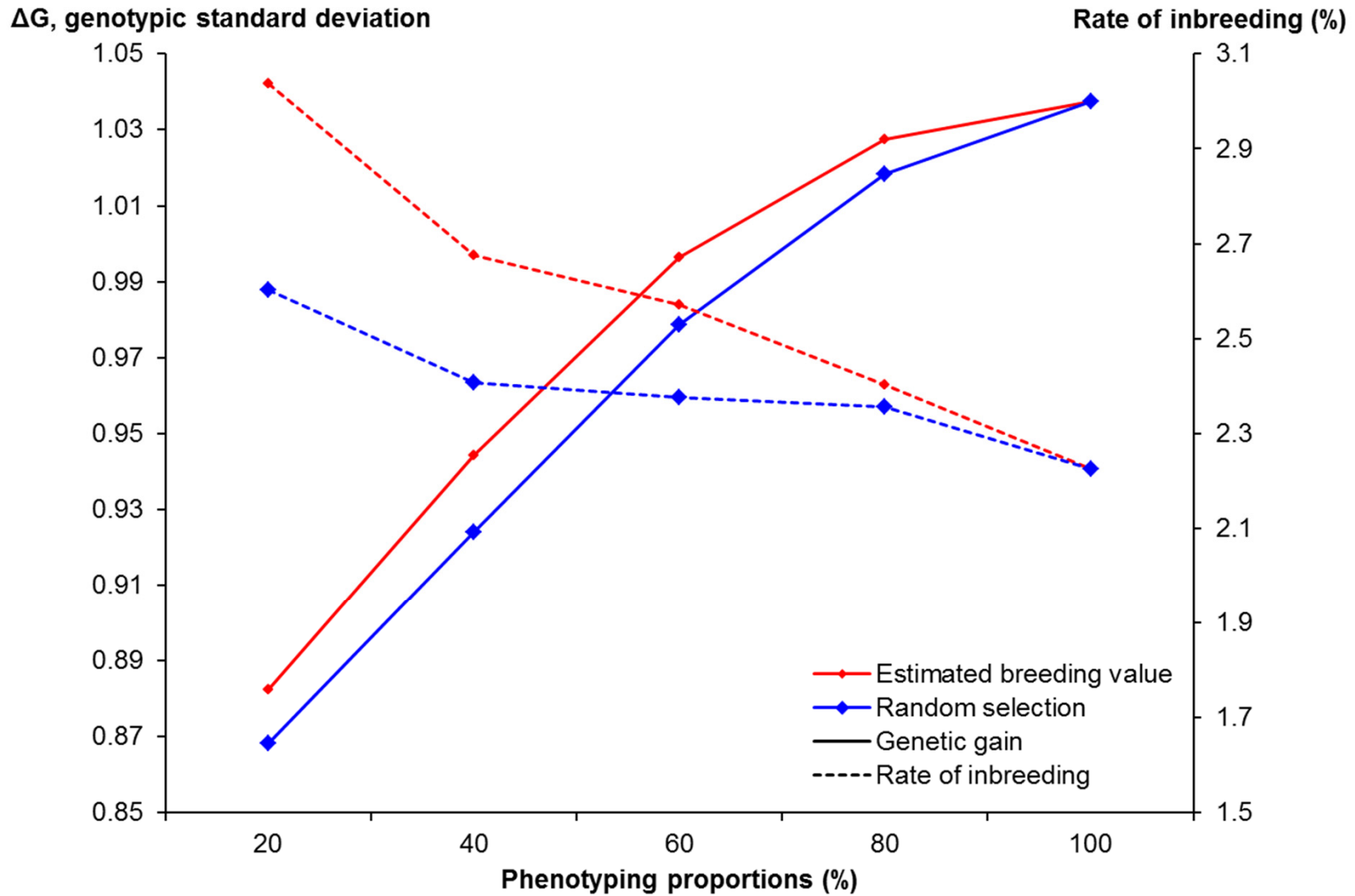
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Previous study - Diminishing marginal returns from genomic selection as more selection candidates are phenotyped- *WCGALP*

➤ **Hypotheses**

- ✓ There is diminishing marginal return from genomic selection as more candidates are phenotyped
- ✓ Phenotyping candidates based on *a priori* information is beneficial



- ✓ Diminishing return to genomic selection as more candidates are phenotyped

- ✓ Use of *a priori* information to select phenotyping candidates is beneficial

- ❑ Most benefits of genomic selection can be realised by phenotyping only high ranking selection candidates

- ❖ The effect of phenotyping candidates' sex ratio on returns to genomic selection was not considered

Objective

- To determine the effect of male-to-female ratio of the phenotyping candidates on returns to genomic selection

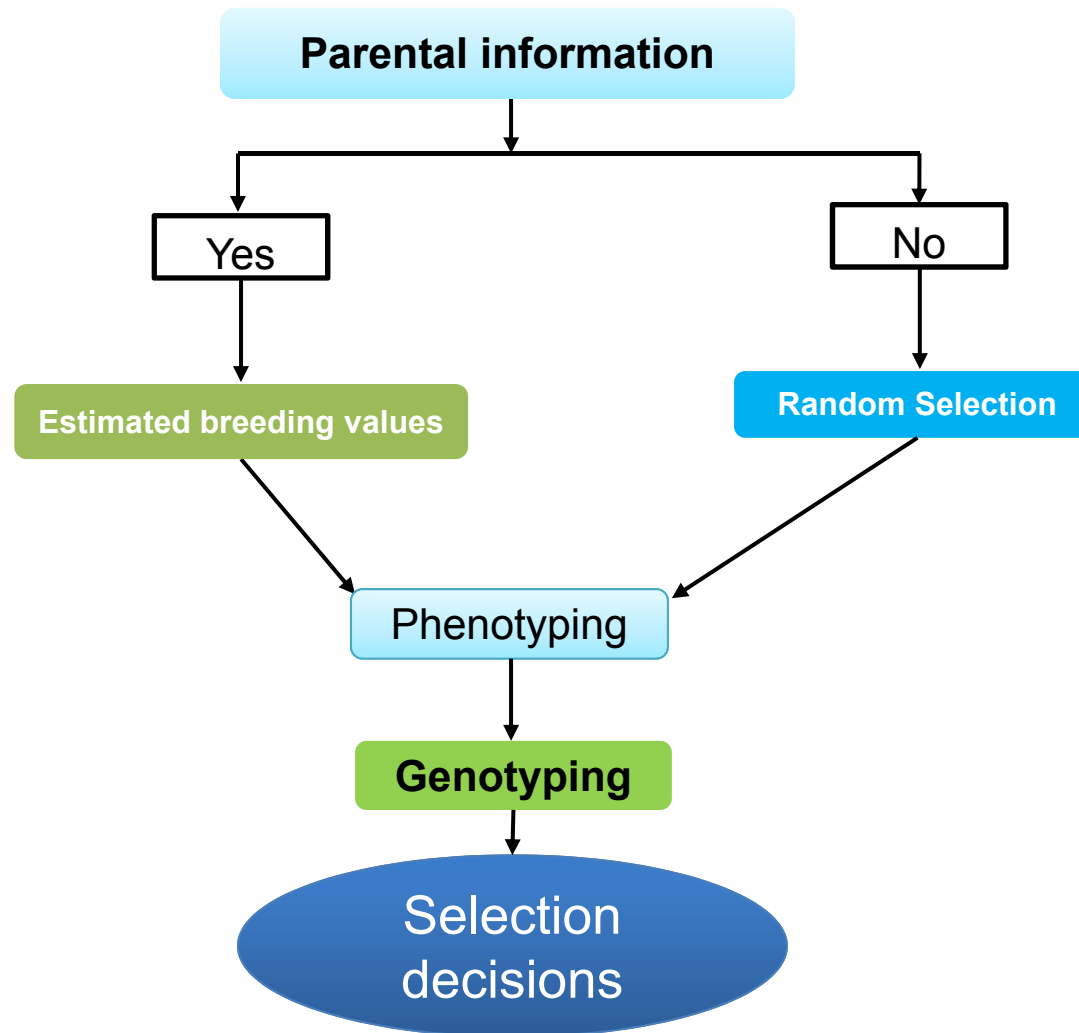
- ❑ **Simulation design**
 - ✓ Genetic gain

- ❑ **Phenotyping proportions**
 - ✓ 20-50%

- ❑ **Male:Female sex ratio**
 - ✓ 100:0
 - ✓ 75:25
 - ✓ 50:50
 - ✓ 25:75
 - ✓ 0:100



□ Phenotyping criteria



□ Breeding scheme

➤ Population structure

- ✓ Boars = 10
- ✓ Sows = 100
- ✓ Litter size = 5

➤ Implementation

- ✓ 12 discrete generations
- ✓ 8-12 - Implementation of genomic selection and phenotyping criteria

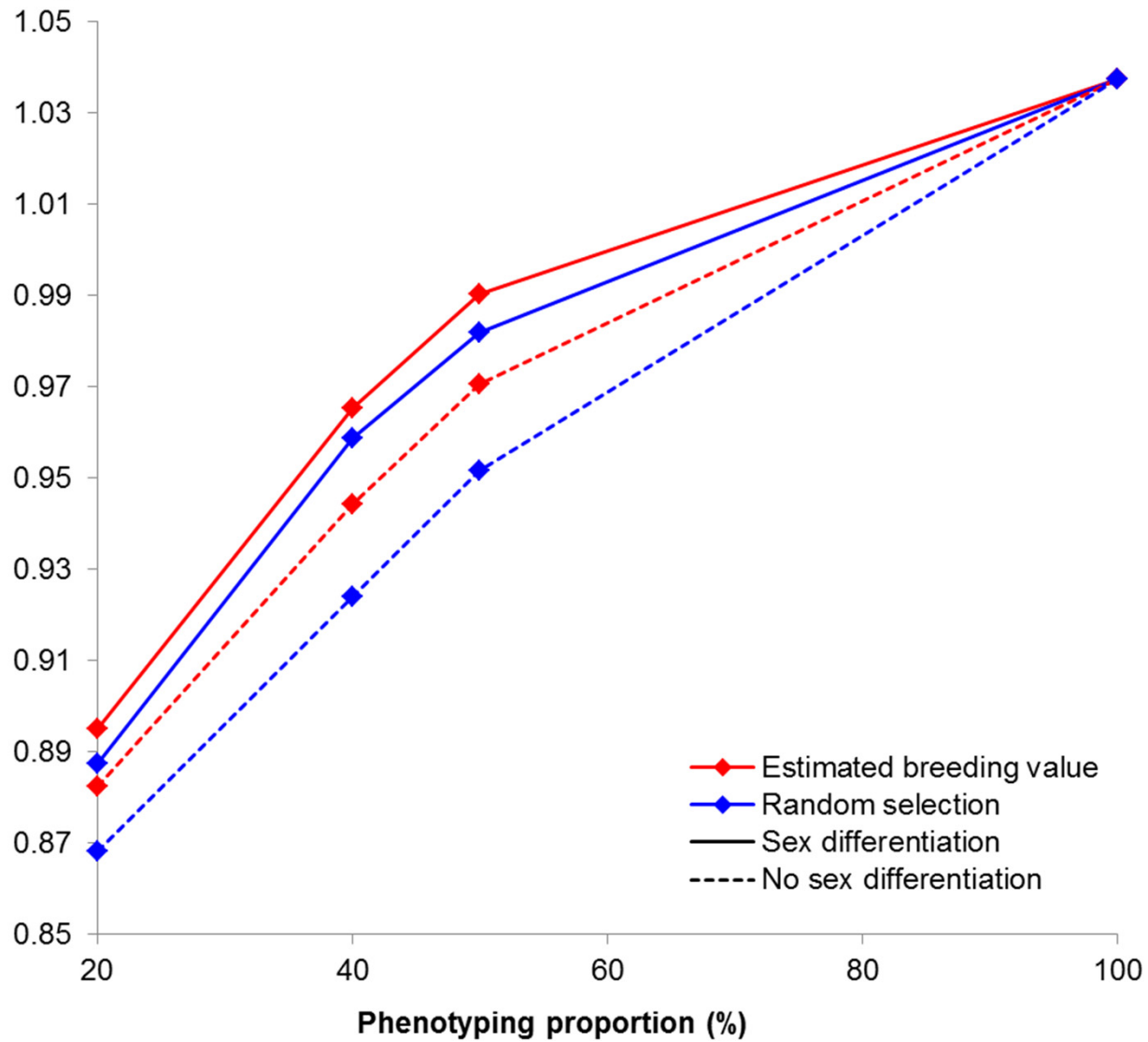
➤ Breeding objective - Single trait selection with $h^2 = 0.4$



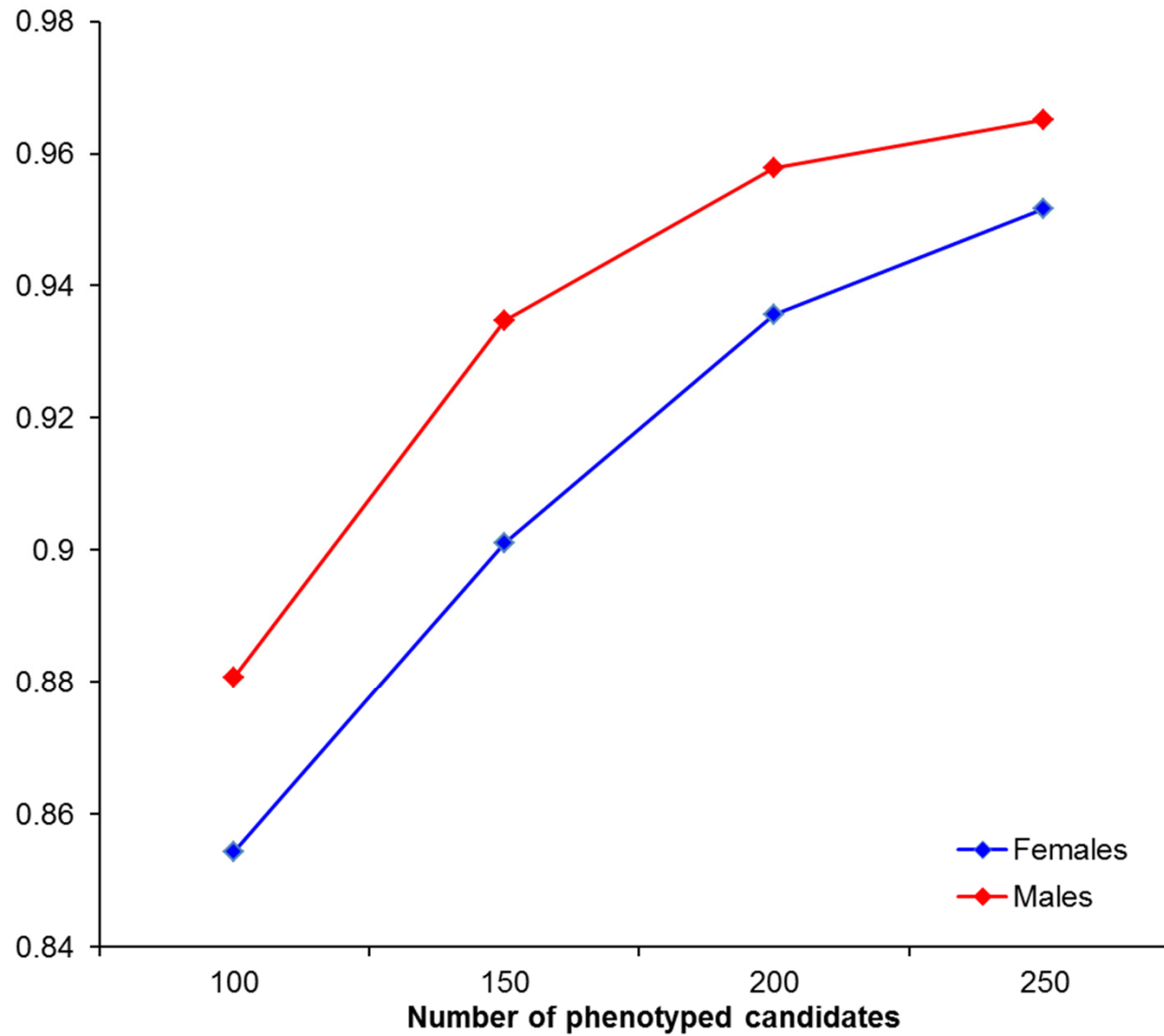
➤ Simulated genome

- ✓ Genetic architecture of the founder population– represent LD in the Danish pigs
- ✓ Genome = 18 chromosomes of 167 cM each
- ✓ 60,000 markers and 8,000 QTLs
- ✓ Haplotypes sampled to initiate breeding schemes

ΔG , genotypic standard deviation



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ΔG , genotypic standard deviation

