

Deriving genotypes of non-genotyped horses

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KWPN



- Studbook Netherlands-based, international oriented
- o Breeding goal: breeding a competition horse that can perform at Grand Prix level
- 20,000 members
- o 11,000 foals per year
- 4 breeding directions:
 - Jumping horses
 - Dressage horses
 - Harness horses
 - Gelders horses









Scope

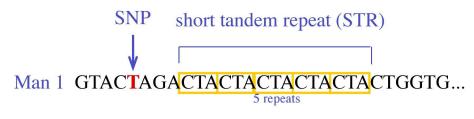


- Genotyping all live-born foals since 2019
 - Neogen: Illumina's Equine 70k SNP chip
 - Parentage verification
 - Breeding value estimation
 - Monogenic abnormalities
- Open population
 - Semen usage from competing studbooks
 - STR genotypes

STR vs SNP



- o STR
 - Repeats
 - 9 mandatory markers
- o SNP
 - More accurate
 - 197 SNPs



Man 2 GTACAAGACTACTACTACTACTACTACTACTGGTG...

Man 3 GTACAAGACTACTACTACTACTACTACTACTACTGGTG... 7 repeats

Research question



What number of genotyped offspring (NOFF), given the percentage of genotyped mates (%MG), is needed to obtain the minimum required accuracy of a derived genotype for use in further applications?

Material and methods

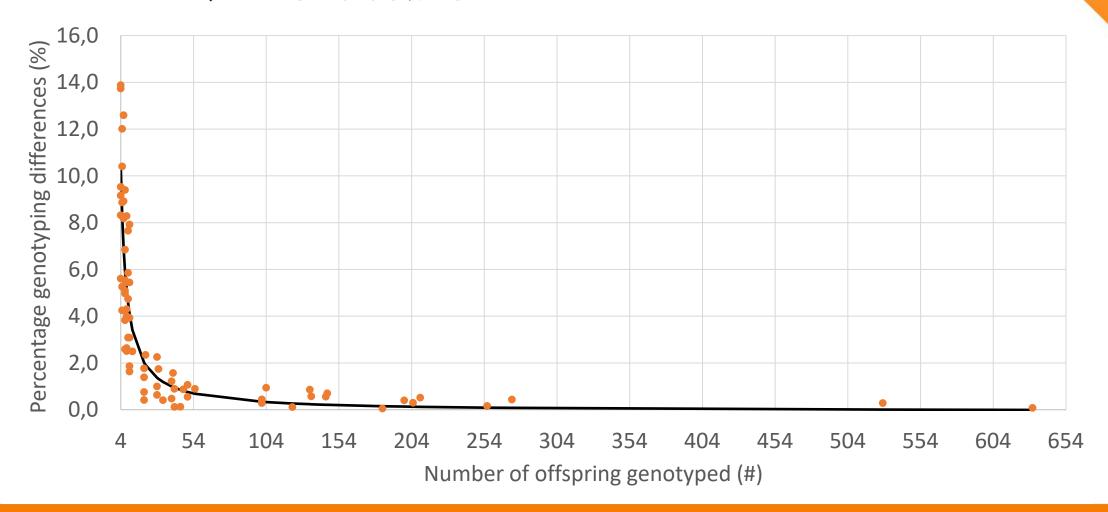


- FImpute (HiggsGene Solutions inc.)
 - Genotypes of sires were removed
 - Derived genotypes compared to observed ones
- \circ 75 sires selected in wide range in terms of NOFF \bar{x} =61 (4..631) and %MG \bar{x} =32 (0..100)
- o Criteria:
 - Call rate >95%
 - Sex chromosomes and the SNPs with a minor allele frequency < 1% were removed
 - 56256 SNPs remained
- Trait of interest: percentage of SNPs that differed between the observed and the imputed ones (%DIFF).
- For establishing the prediction equation for %DIFF, a multiple regression analysis was performed where the reciprocal of NOFF and its interaction with %MG were used as independent variables.

Results

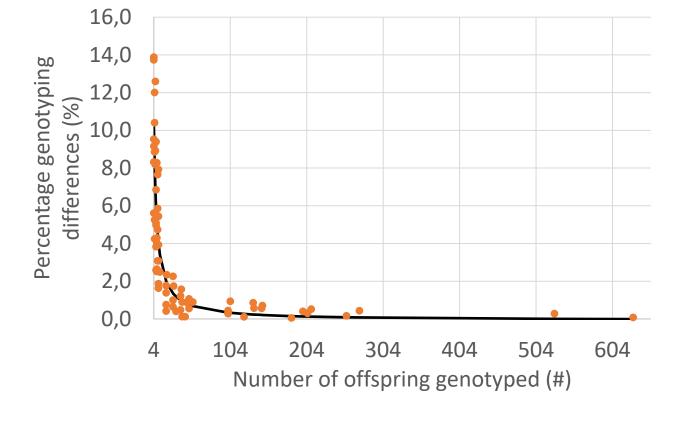


○ %DIFF = $0 \rightarrow 50$ NOFF and 100 %MG $\rightarrow 127$ NOFF and 0 %MG



Results

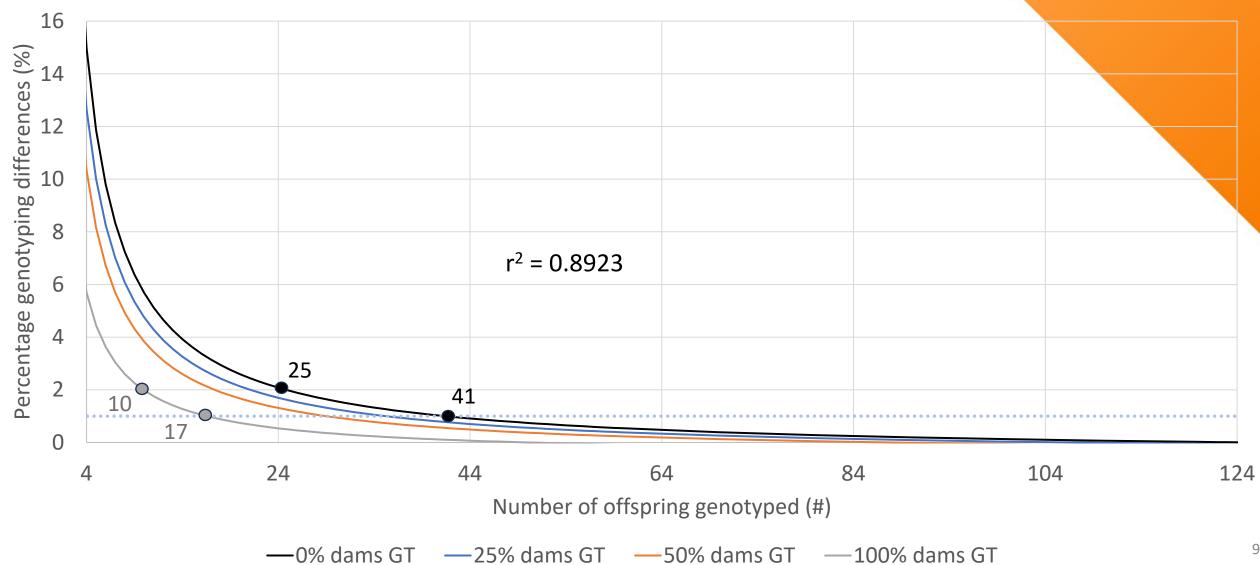
- %DIFF = $0 \rightarrow 50$ NOFF and 100 %MG $\rightarrow 127$ NOFF and 0 %MG
- The r² of the prediction equation was 0.89
- Both independent variables were significant (p <0.0001).



- Accuracy of derived genotypes is bigger than for observed genotypes.
- %DIFF = 2 \rightarrow 10 NOFF and 100 %MG \rightarrow 25 NOFF and 0 %MG

Results





Into practice



- Sire only has STR profile
 - ~1100 foals per year are genotyped for both STR and SNP
 - Double costs
- Create routine to impute missing genotypes

Opportunities



- Parentage verification KFPS (Frisian horses)
 - Started end 2023
 - STR is needed
 - Colour traits, Hydrocephalus and Dwarfism
- Lowers the bar to switch to SNP genotyping for other studbooks



Thank you for your attention

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