



# First Steps towards Genomic Selection in French Dairy Goat

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# Few males progeny tested each year in French dairy goats

Progeny tested bucks each year

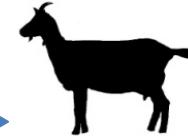
40 Alpine



30 Saanen



100 daughters /year



with performances

25 Alpine

AI bucks

15 Saanen

CD<sub>asc</sub> : 38%



> 5.5 years  
sire-son  
pathway



Alpine or Saanen

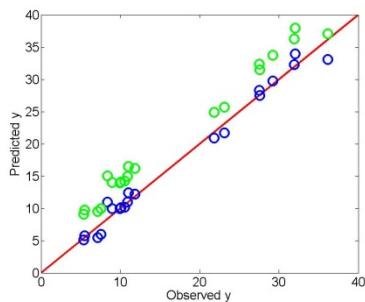


2011: Illumina goat SNP50 Bead Chip

# Not ideal population structure: prediction reliability similar to dairy sheep



**linkage disequilibrium**



**genomic evaluation**



**interest of adding females in reference population**

# Small multi-breed population genotyped with females

46 959 SNP after quality control



384 Alpines



293 Saanens

QTL  
detection



1,243 Alpines

1,985 females



742 Saanens



87 Alpines

148 males

not progeny tested yet



61 Saanens

1993

2008

2009

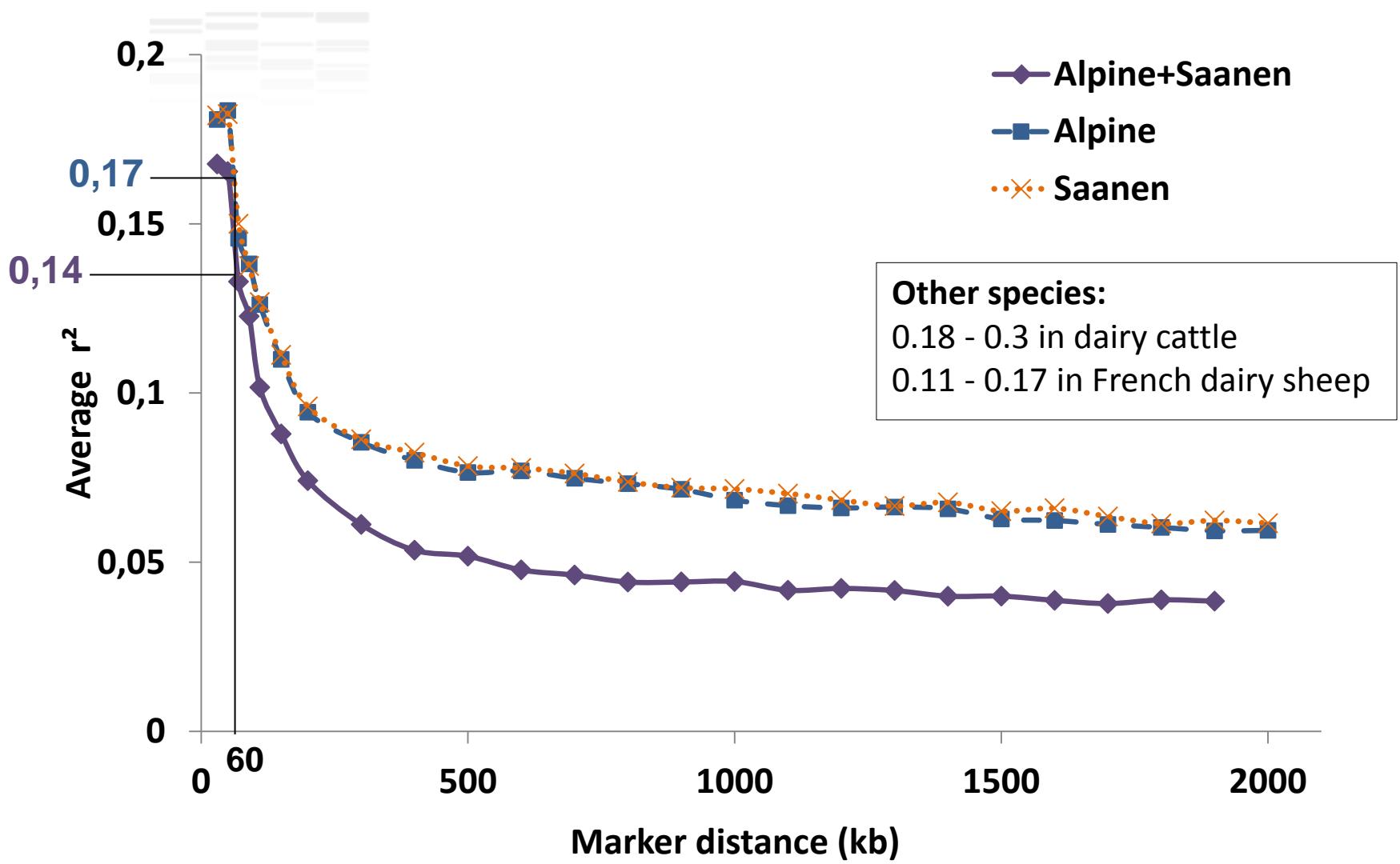
2010

2011

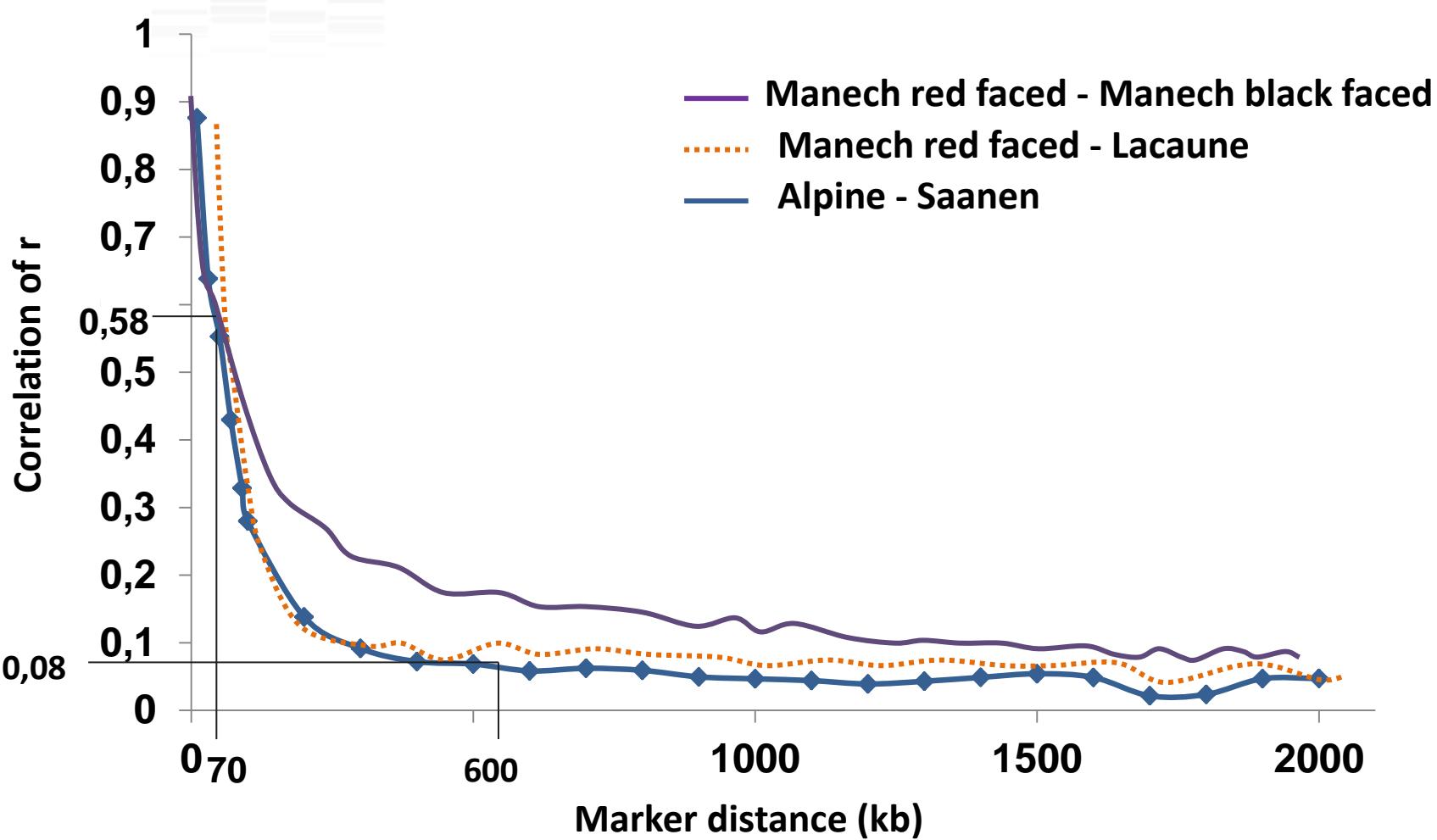


.04

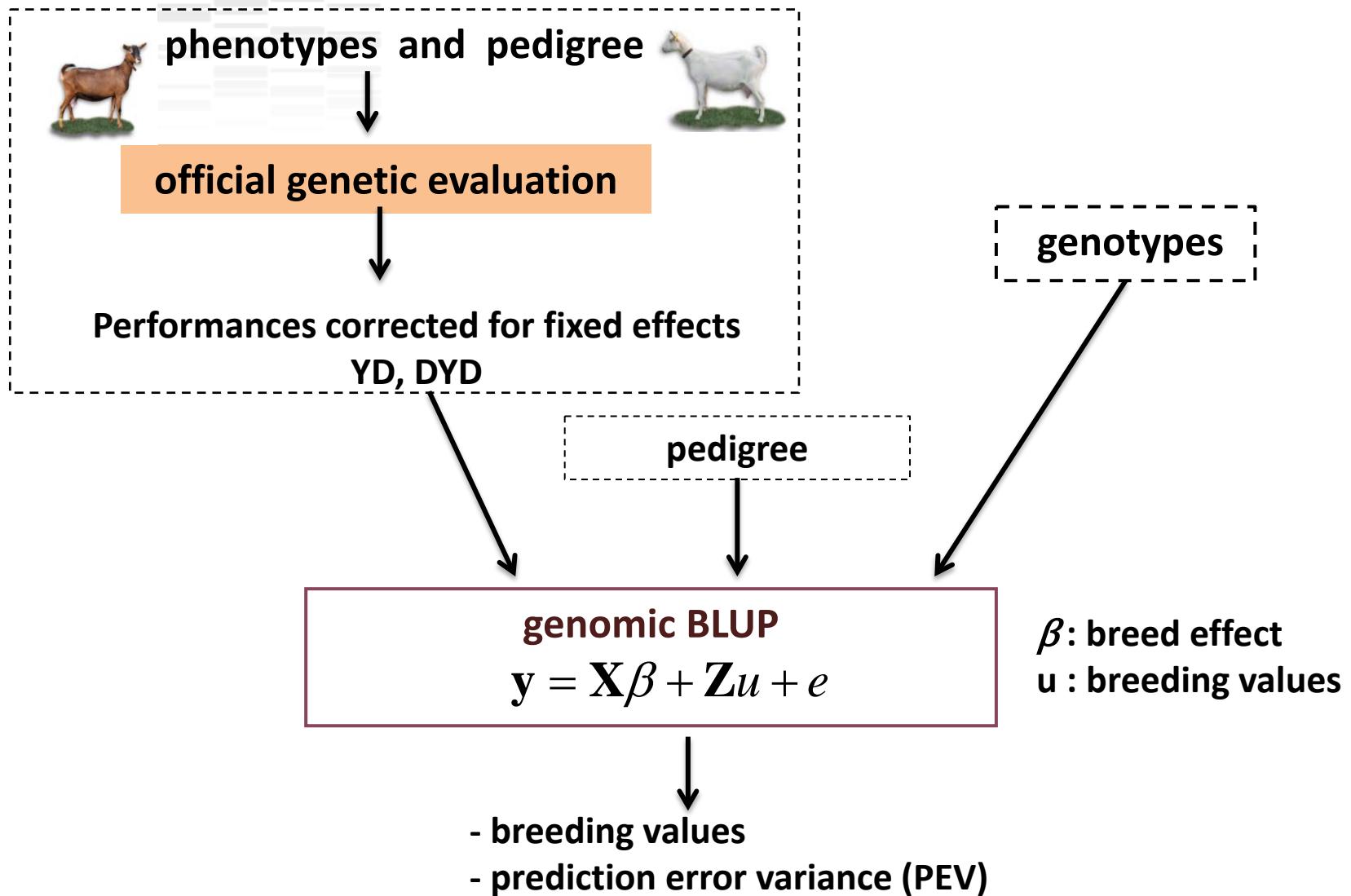
# Medium linkage disequilibrium in dairy goats, lower in multi-breed population than in each breed



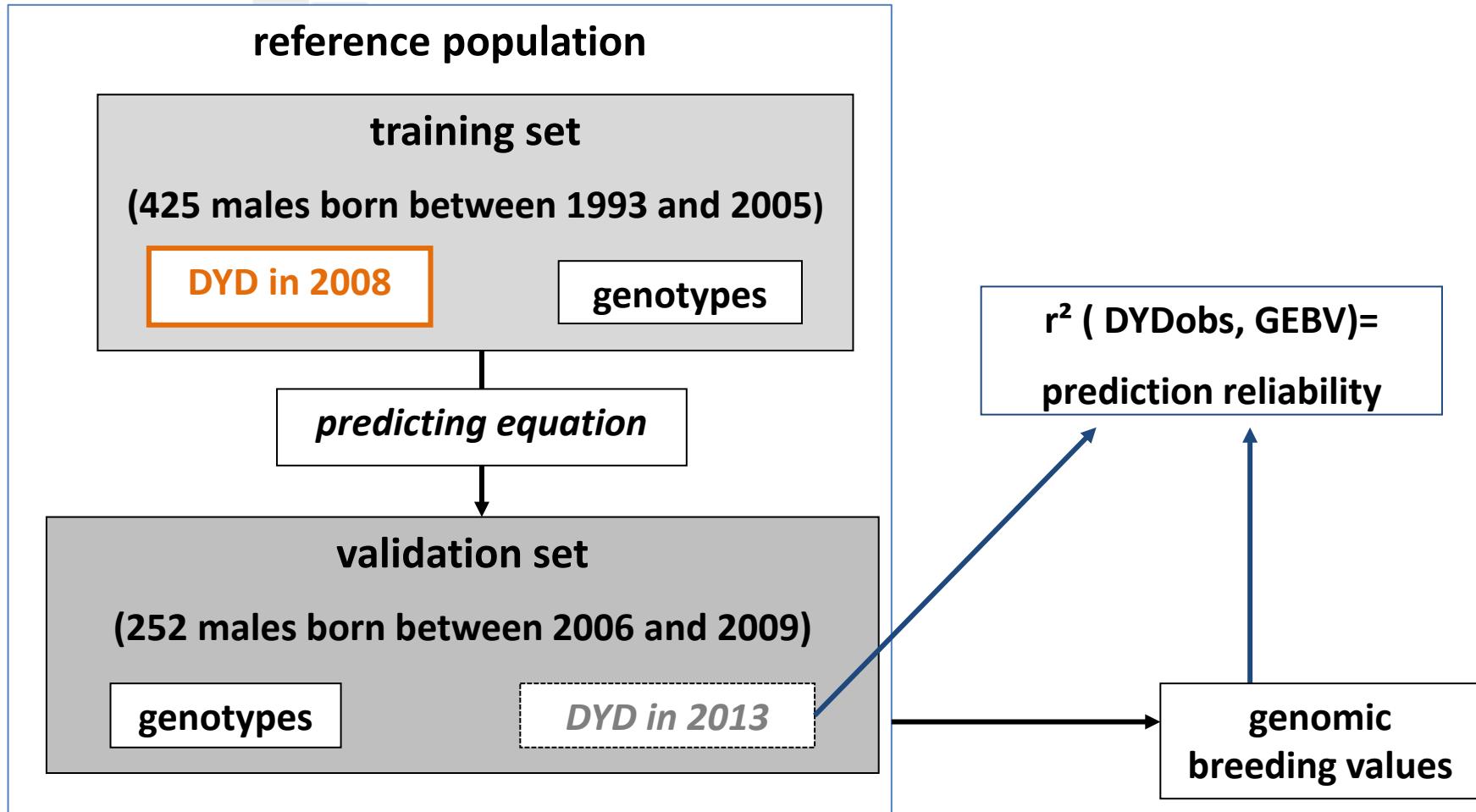
# Alpine and Saanen : two different breeds according to LD persistence between both



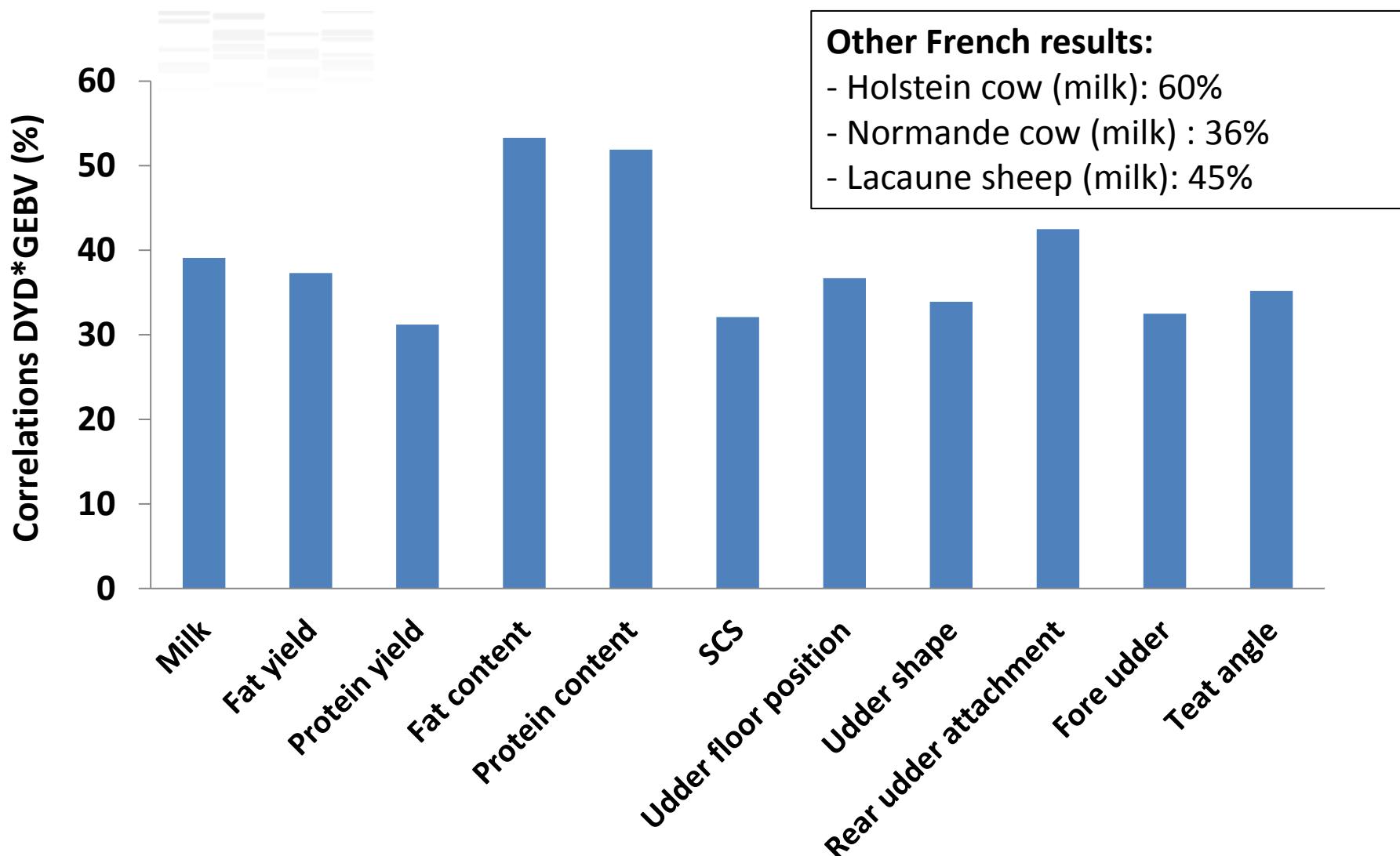
# What are the model and the data used in this study?



# How we calculate prediction reliability?



# Prediction reliability in validation population close to the one in Normande dairy cattle breed



# No males with daughters born after females genotyped

46 959 SNP after quality control



677 males

384 Alpines



293 Saanens

QTL  
detection



1,985 females

1,243 Alpines



742 Saanens



148 males  
not progeny tested yet

87 Alpines



61 Saanens

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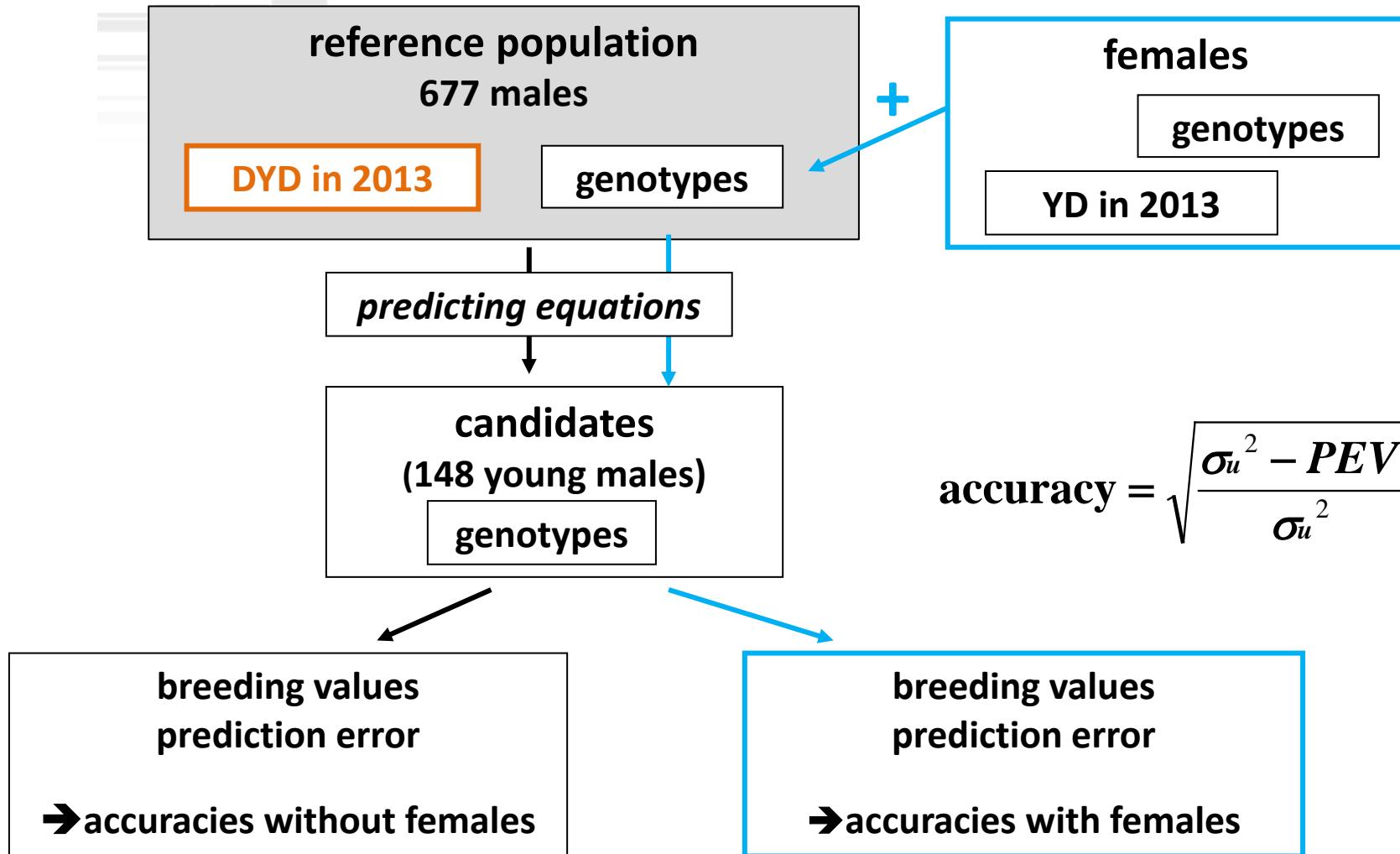
2010

2011



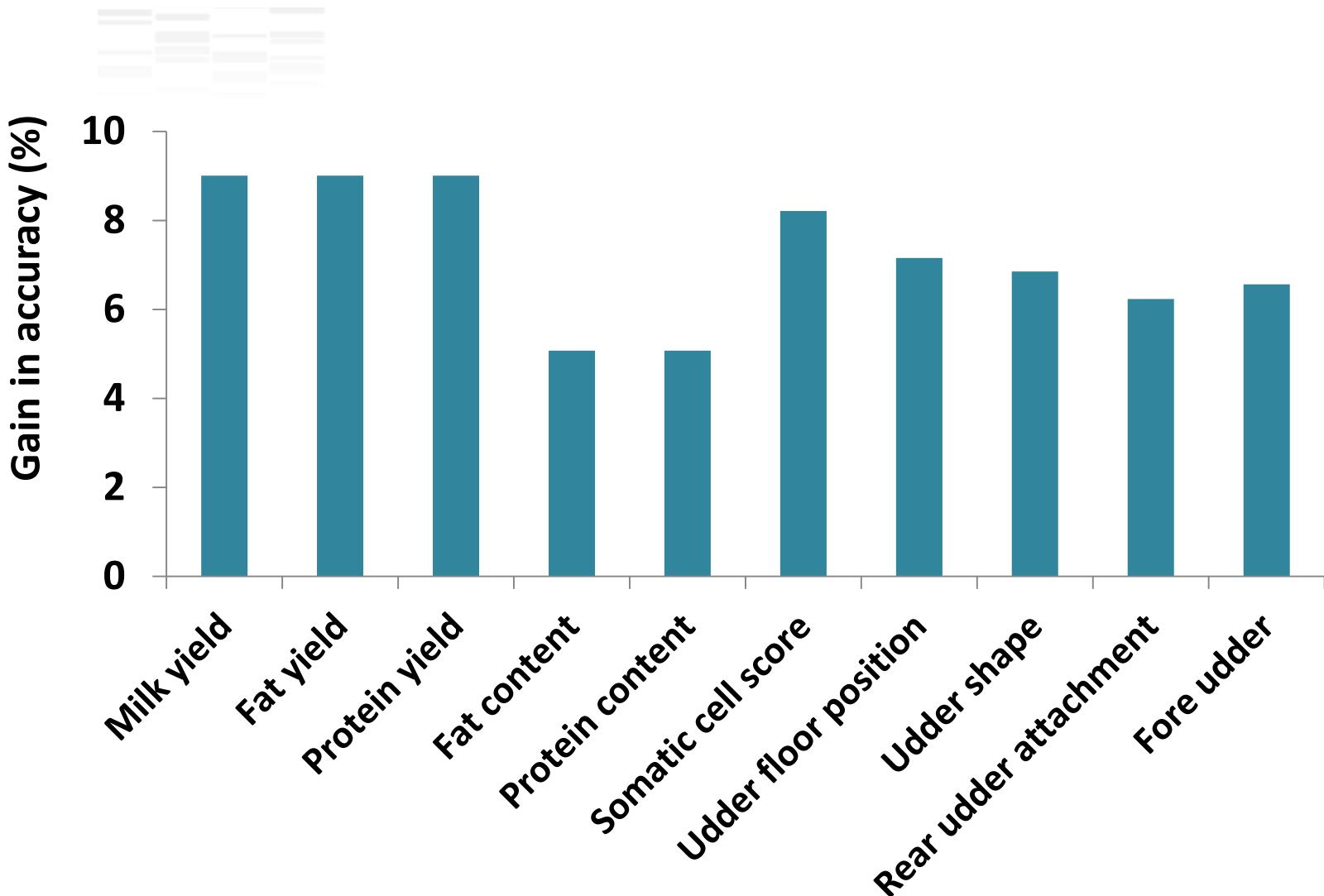
.010

# How we calculate accuracy of young buck breeding values?

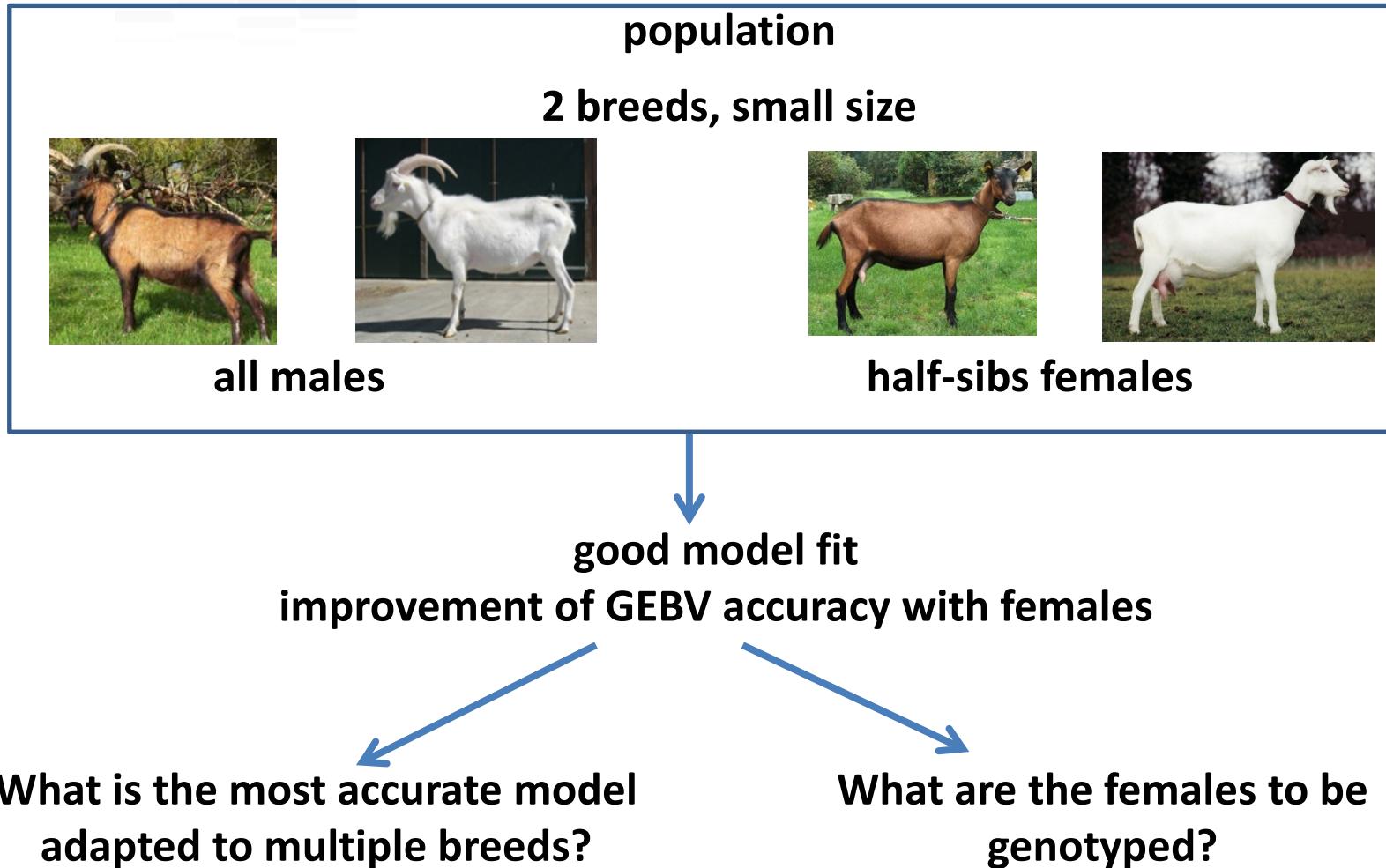


$$gain = \frac{accuracy_{with\ females} - accuracy_{without\ females}}{accuracy_{without\ females}}$$

# Improvement of accuracies when adding 1,985 females in reference population of 677 males



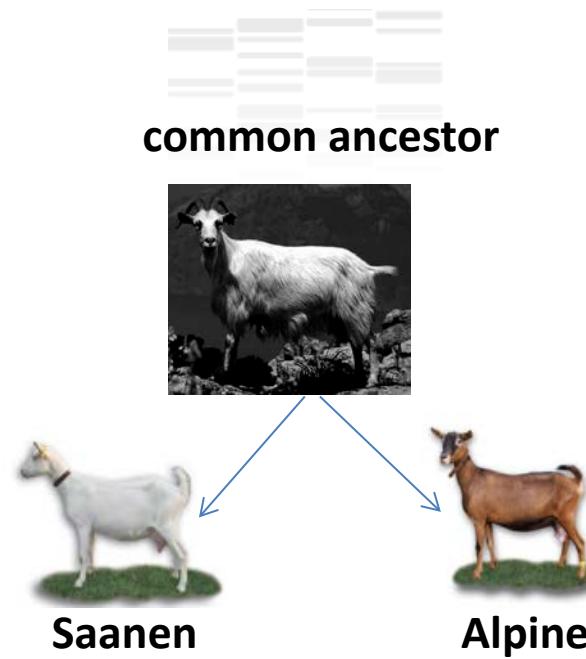
**Genomic selection is difficult to apply due to small population size, two breeds and population structure**





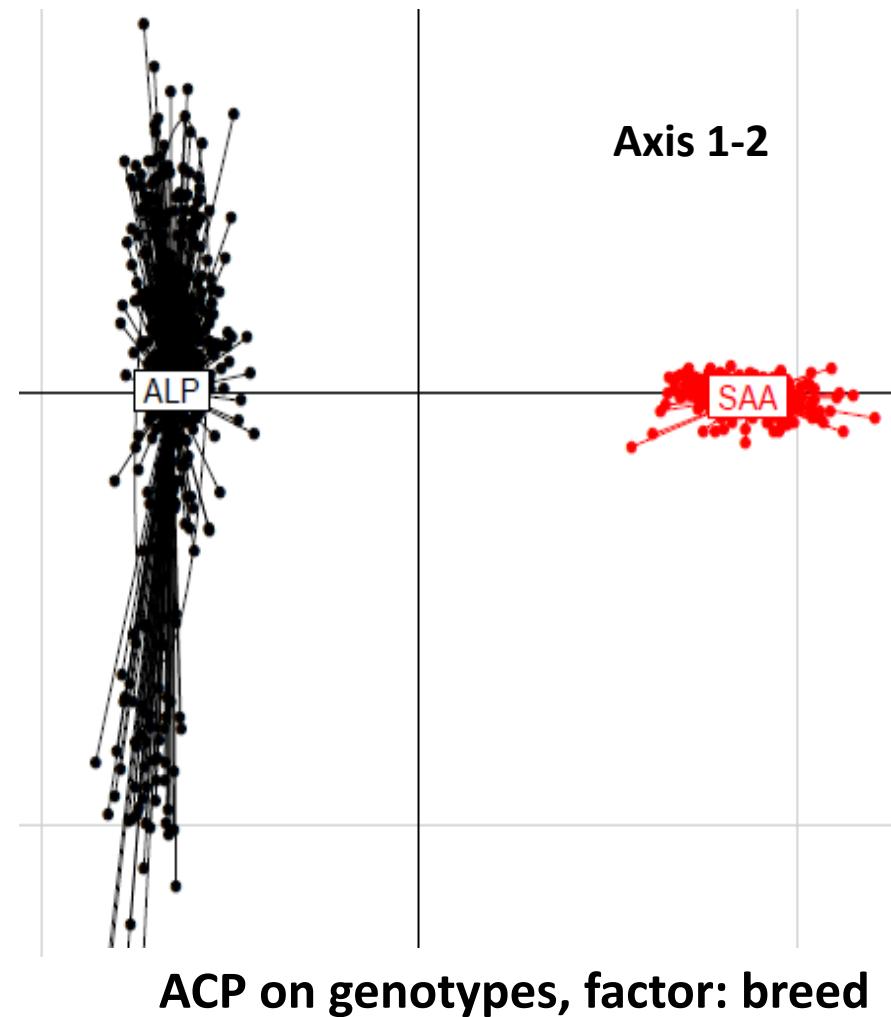
# Appendix

# Difficulties to consider Alpine and Saanen as a single breed



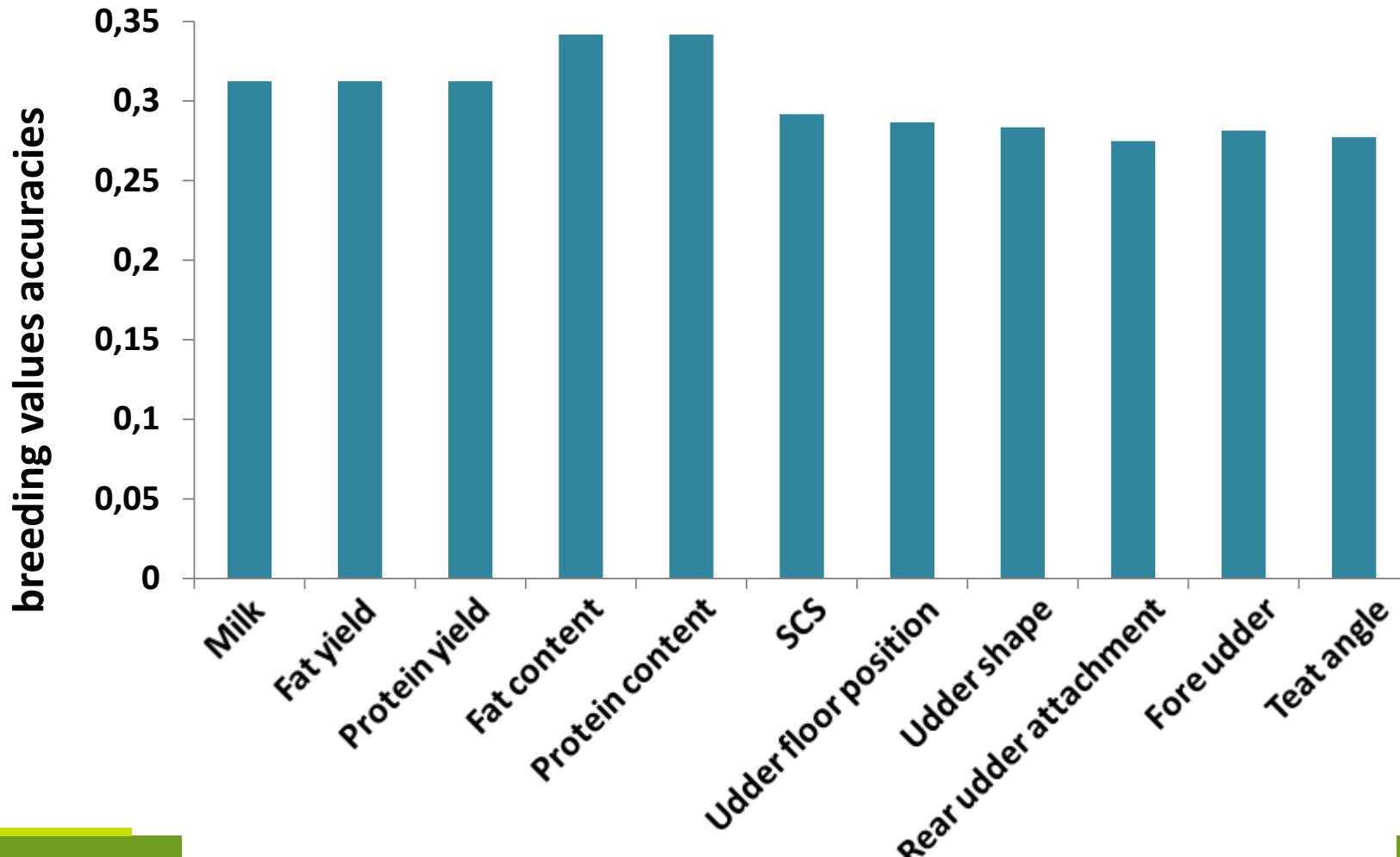
correlation between allele frequencies :

Males	0.6852
Females	0.8455
Both	0.8605

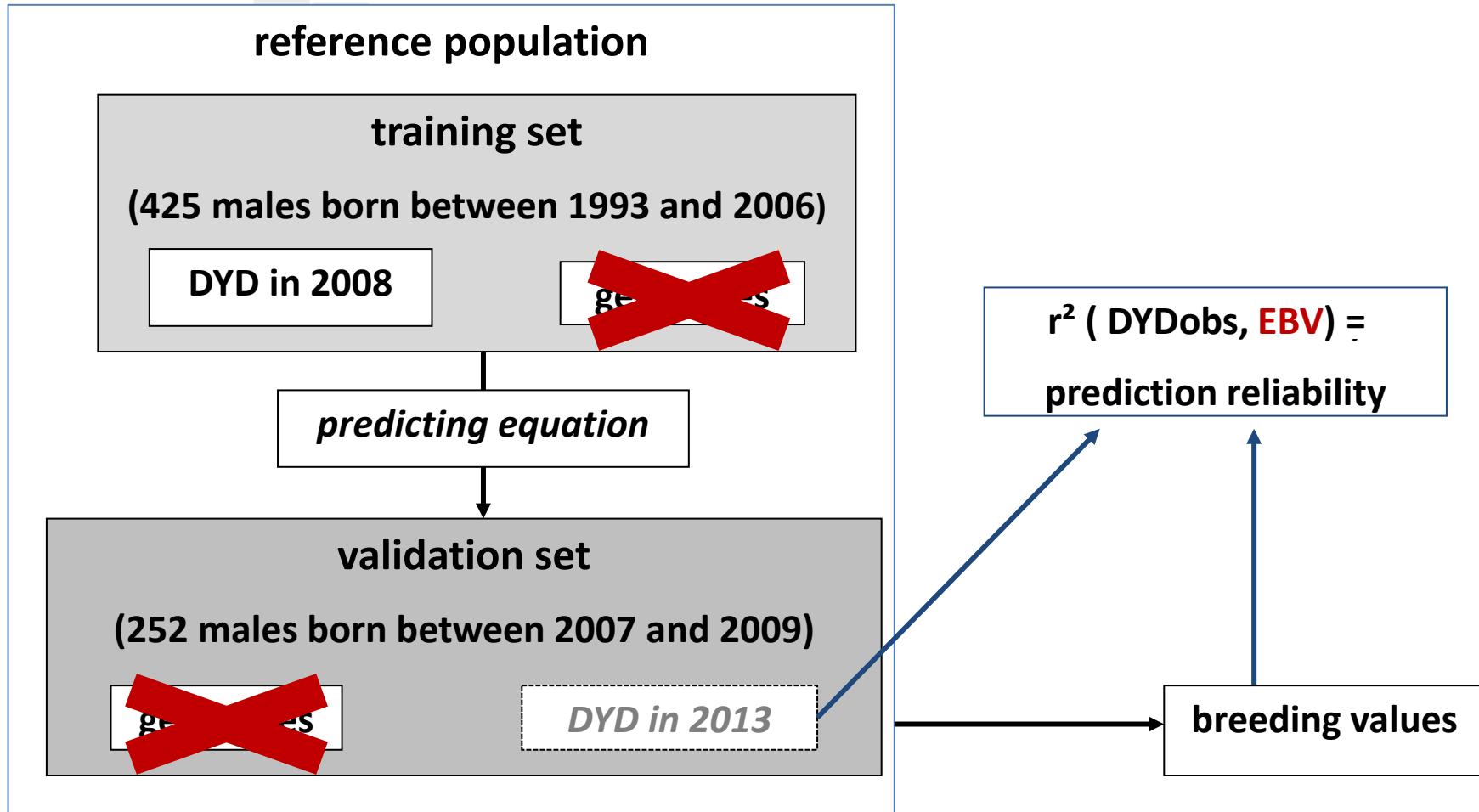


# Young buck breeding value accuracies not as high as expected

Reference population: 677 males and 1,985 females

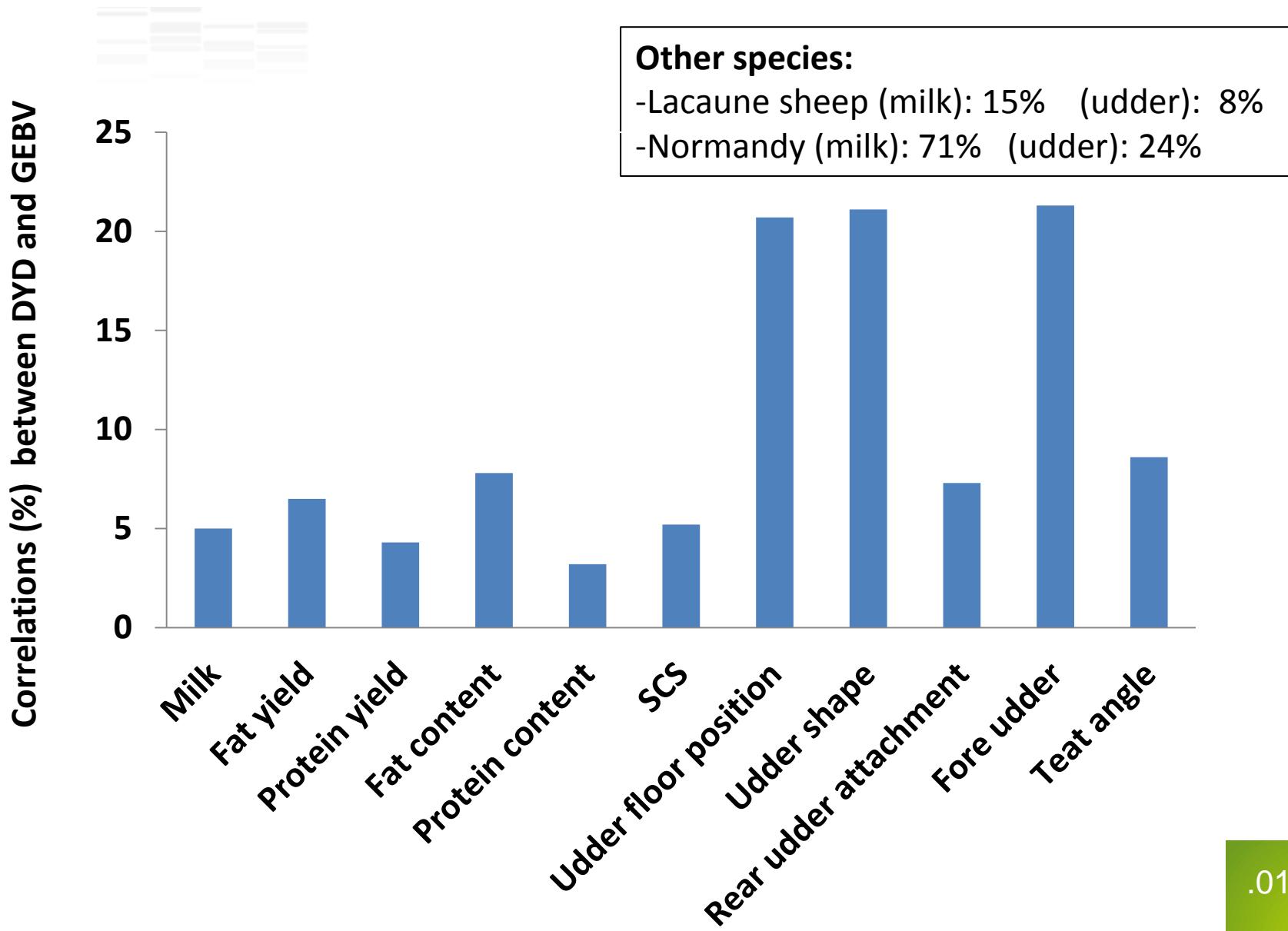


# How we calculate prediction reliability?



$$\text{Gain in reliability} = \frac{r^2(DYD_{obs}, GEBV) - r^2(DYD_{obs}, EBV)}{r^2(DYD_{obs}, EBV)}$$

# Gain in reliability with genomic information lower than in other species for milk production traits



# Few male progeny tested each year in French dairy goats

Progeny tested buck each year

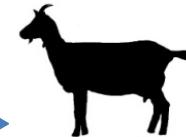
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Sire-son



< 4 years  
Sire-daughter



Alpine or Saanen



2011: Illumina goat SNP50 Bead Chip

# Alpine and Saanen : two different breeds according to LD persistence between both

