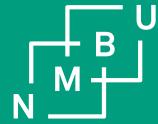


Norwegian University  
of Life Sciences



# Assessing the Genetic diversity conserved in the Norwegian live poultry genebank

Brekke C., Groeneveld L. F., Meuwissen T. H. E., Sæther N., Weigend S., Berg P.

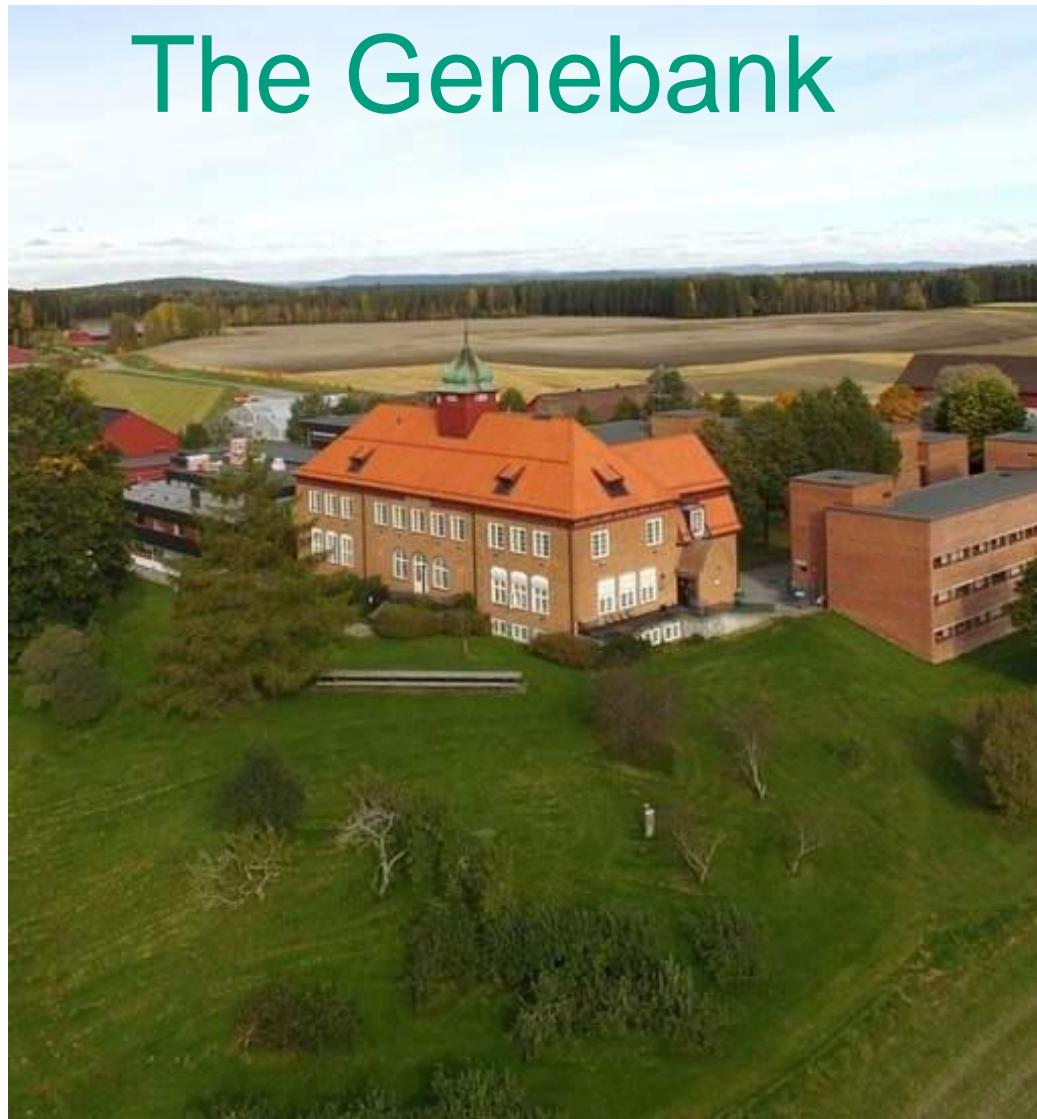
Submitted: *Acta Agriculturae Scandinavica A Animal Science*

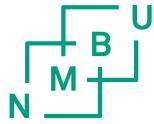


EAAP “What to conserve” 26.08.2019



# The Genebank



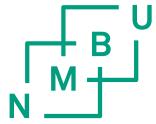


# Lines in the study

Jærhøns - The national breed



19 Individuals genotyped



# Lines in the study

## Previous commercial white layers



# Lines in the study

## Previous commercial white layers



# Lines in the study

## Previous commercial brown layers

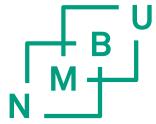




# Lines in the study

Brown eggayers



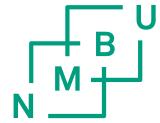


# Lines in the study

## Brown egg layers

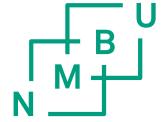


# International reference data



70 different lines from the Synbreed project (Malomane et al. 2019).

# Commercial reference data



Lohmann White



Lohmann Brown



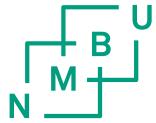
# Genotyping

- **Affymetrix® Axiom® Chicken Genotyping Array**
- High density SNP Array
- 580 000 SNPs



# Aim of the study

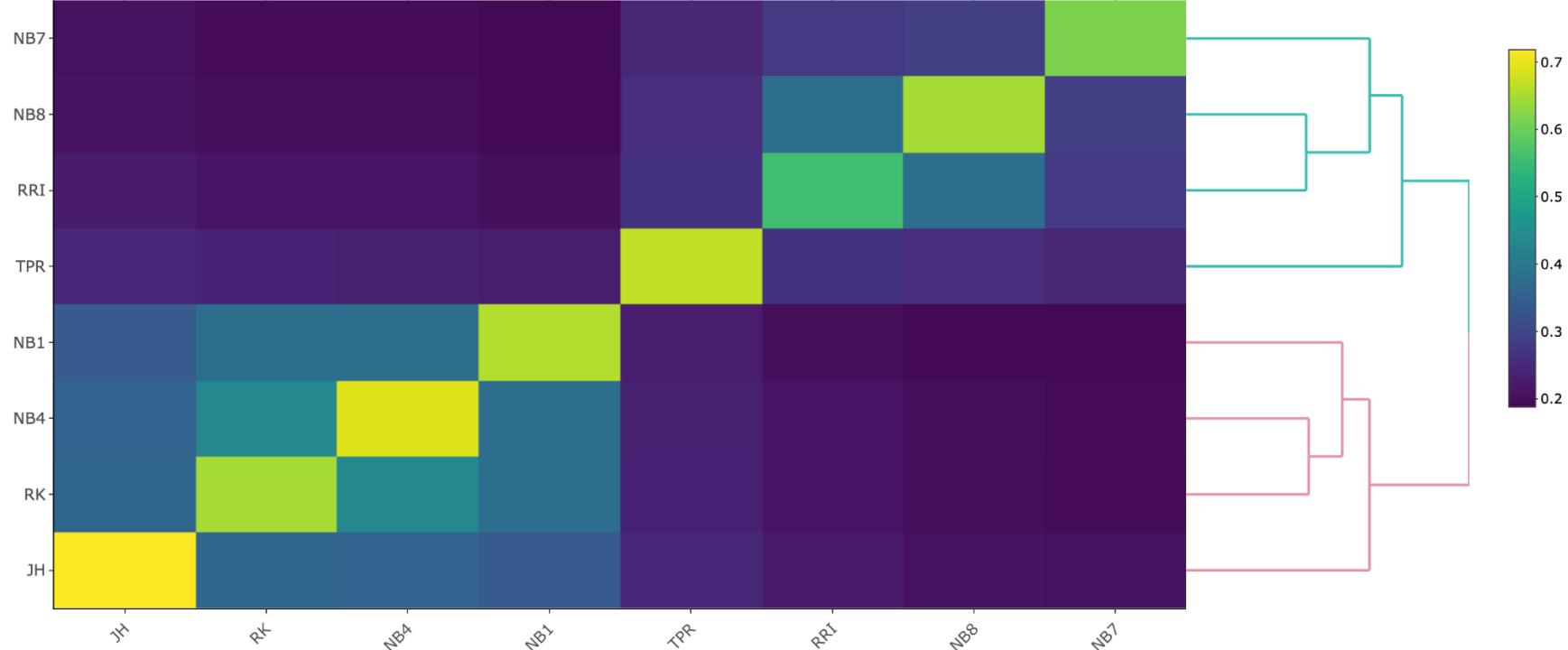
- Estimate genetic diversity within and between the lines
- Quantify conservation value



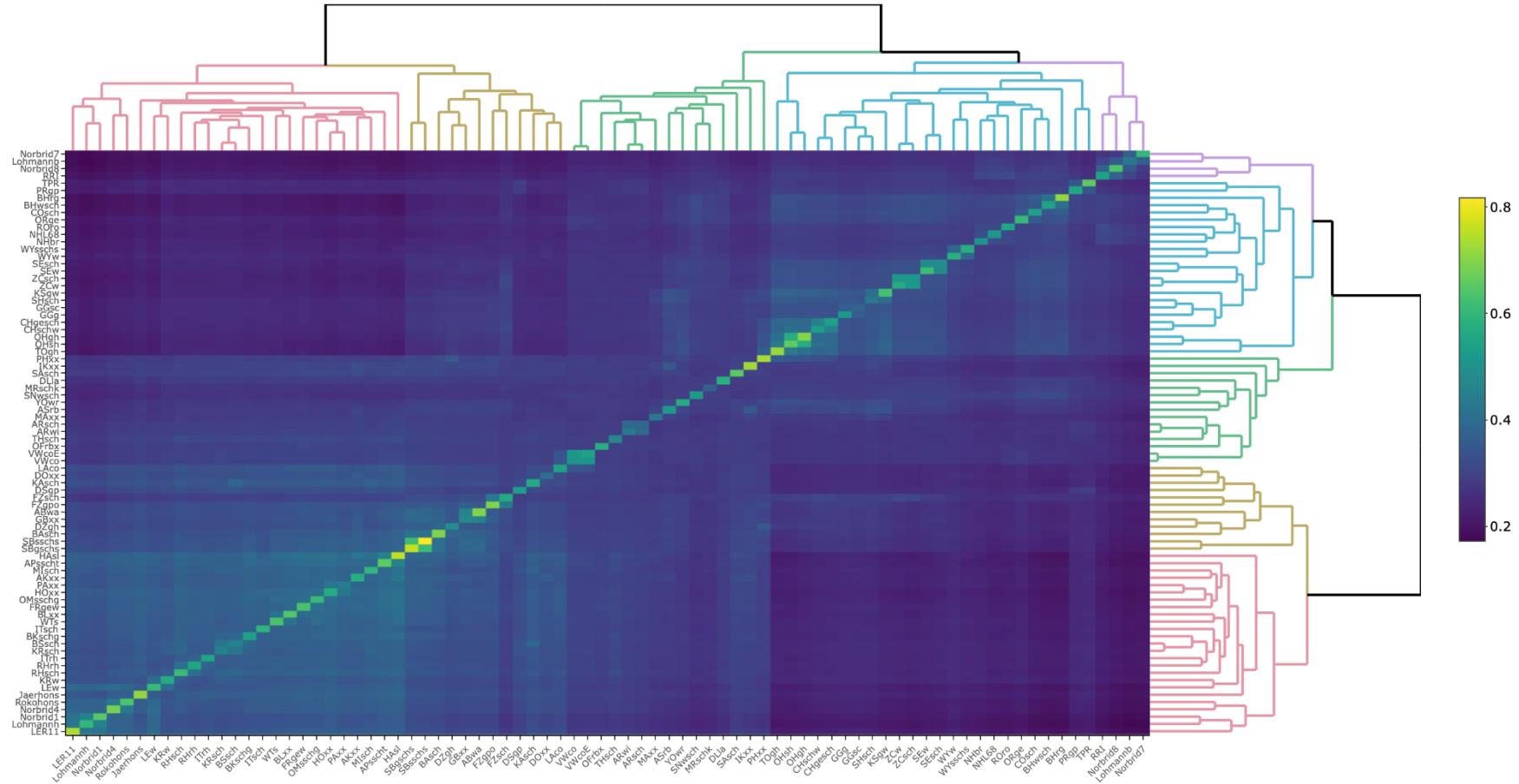
# Average genomic relationships Norwegian lines

$$\mathbf{G} = \frac{\mathbf{Z}\mathbf{Z}'}{2 \sum p_i(1-p_i)}$$

(VanRaden 2008)

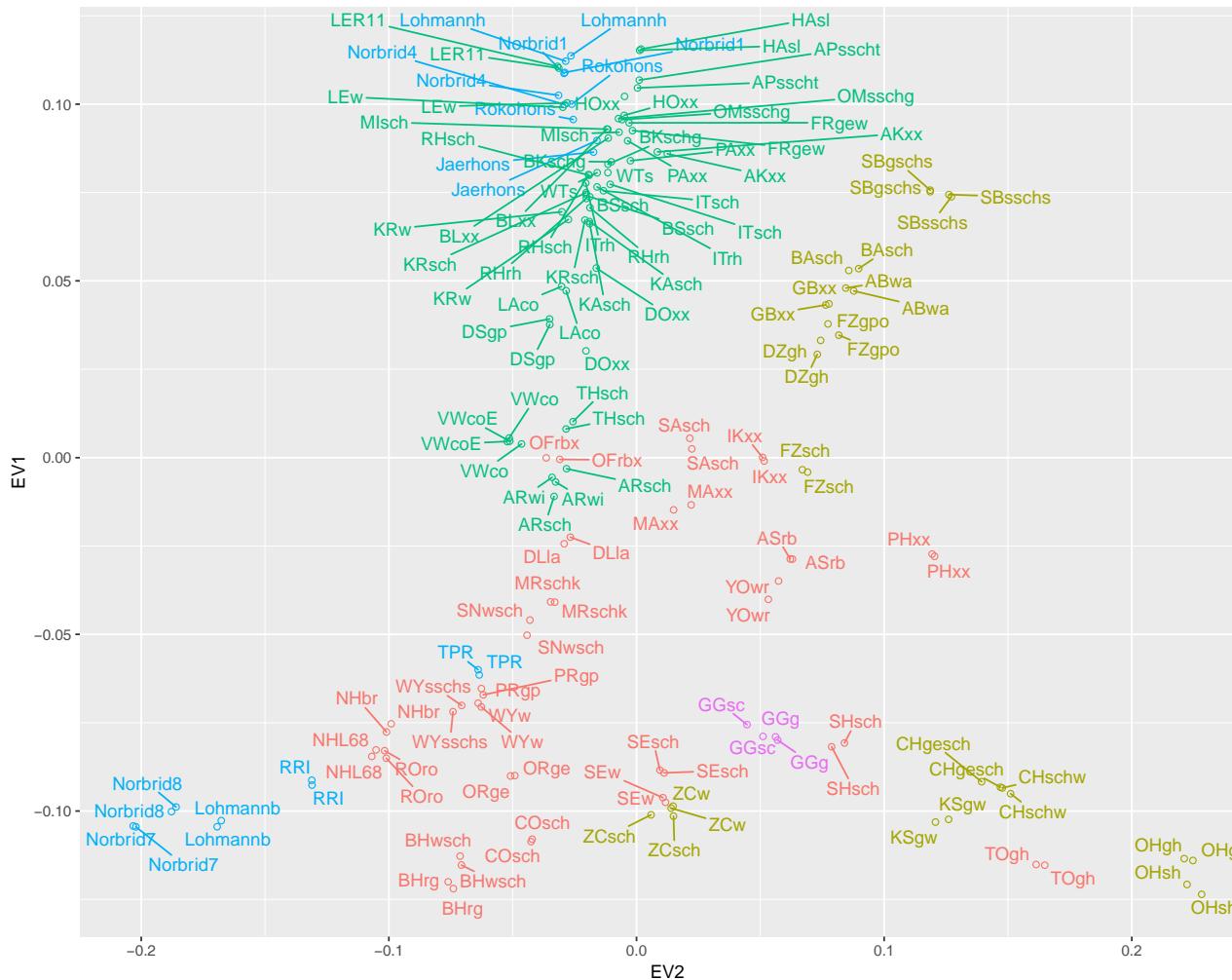


# Average genomic relationships



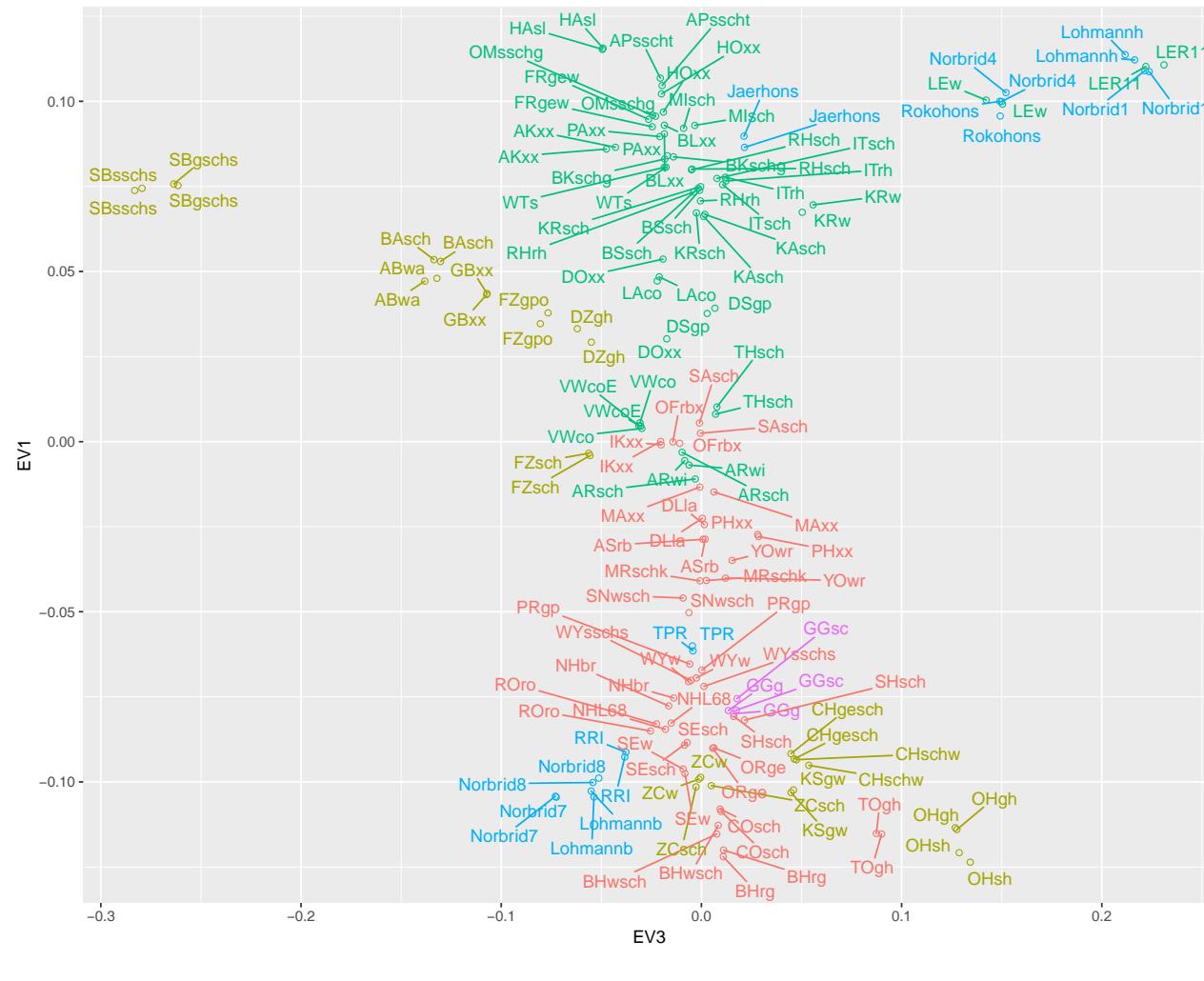
# PCA

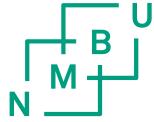
U  
B  
M  
N



# PCA

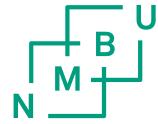
U  
B  
M  
N



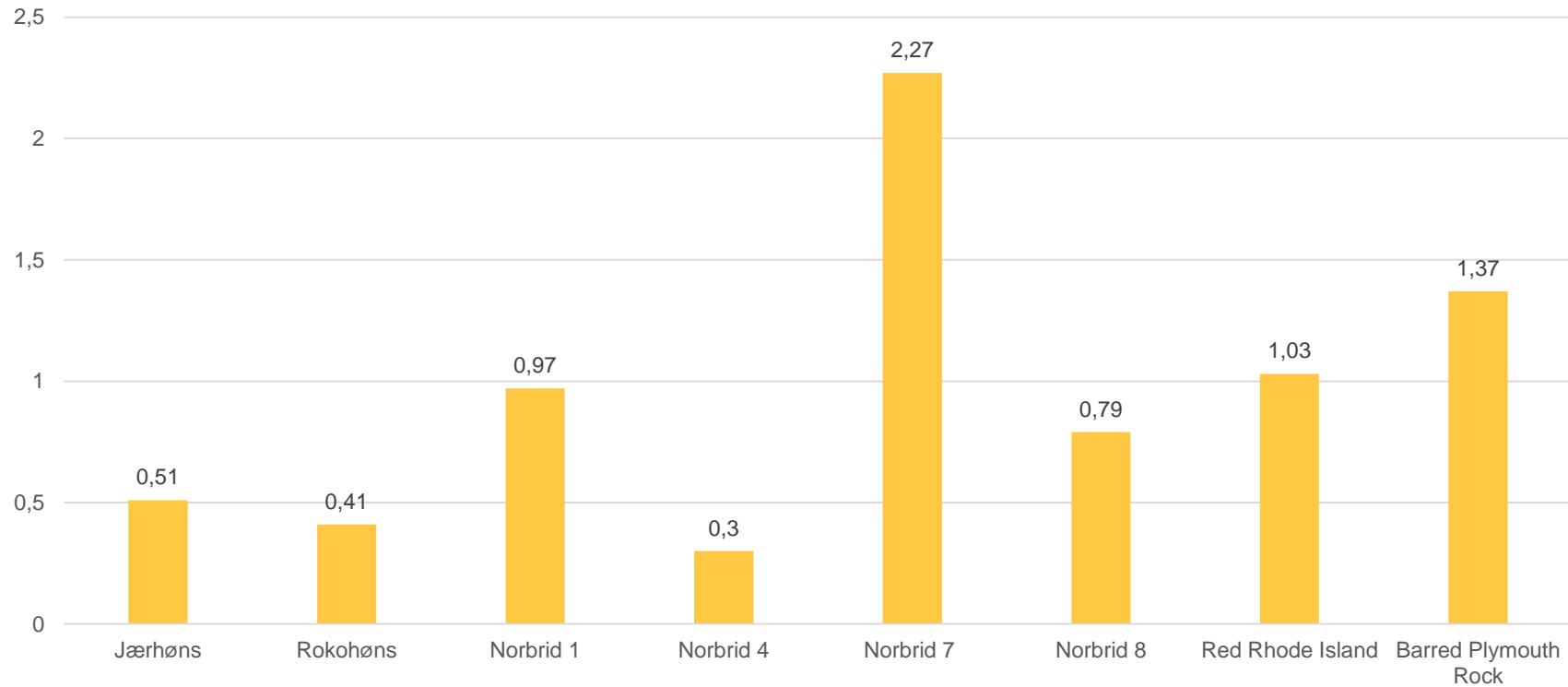


# Contribution to genetic diversity

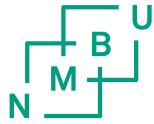
- Core set method suggested by Edings et al. 2002
- Set of breeds is constructed (safe set)
- If adding one breed to the set leads to a decrease in genomic relationships the breed has a contribution to genetic diversity



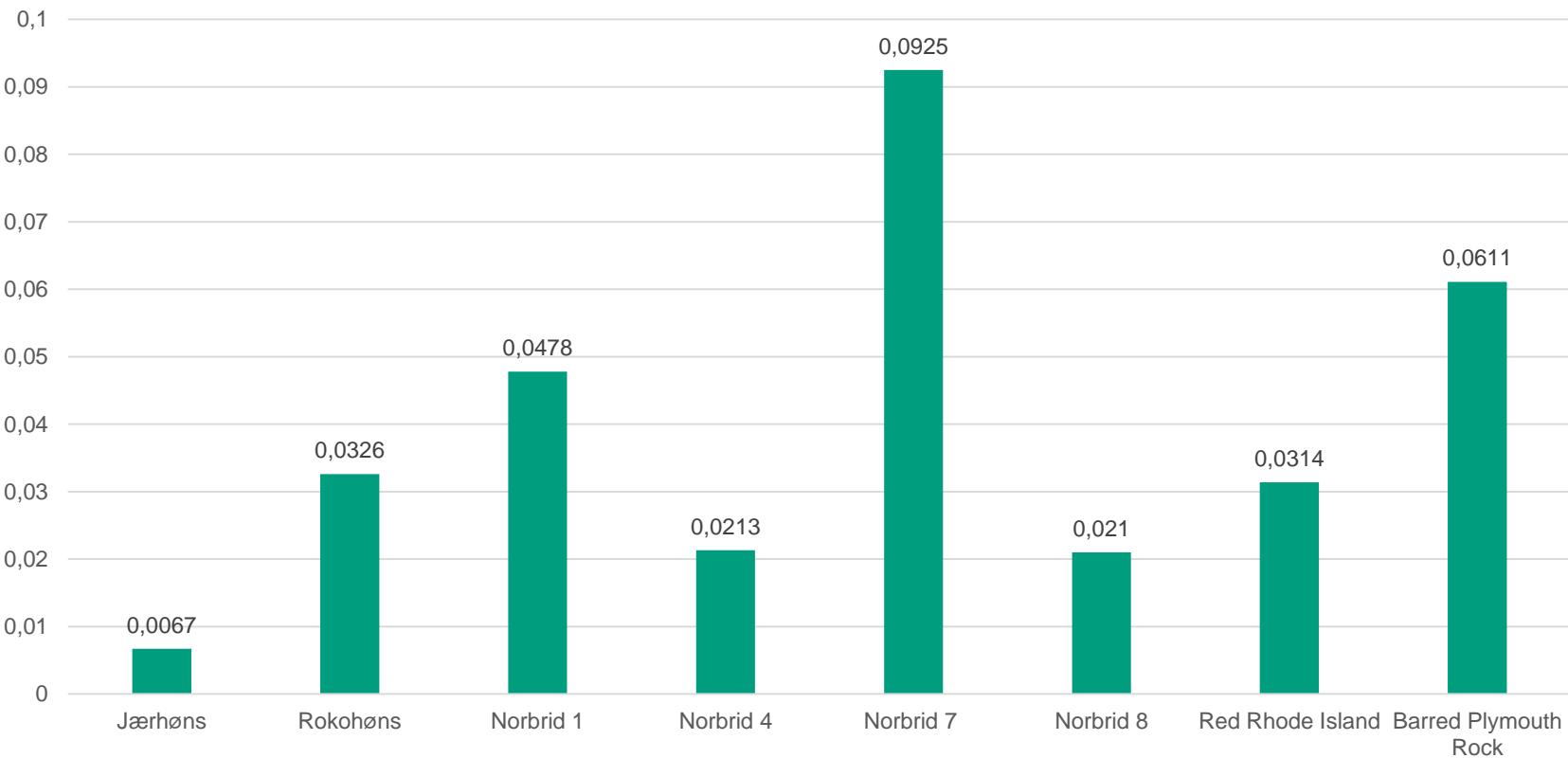
Contributions to genetic diversity (%)  
Core set = Norwegian genebank lines



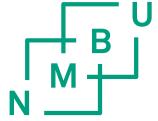
Core set method (Edings et al. 2002)



### Contribution to genetic diversity (%) Core set = International dataset



Core set method (Edings et al. 2002)



# Discussion

- Diversity across the genome, or in specific regions or for specific traits of interest?
- Phenotypic diversity does not necessarily translate into genomic diversity (and vice versa)
- (Previous) Commercial lines = genetic resources



Thank you!