



Introduction

- Valuable recording new traits in research herds
 feed intake, progesterone for fertility, disease resistance ...
- Any use pooling data across research herds for genomic selection?

ROBUSTMILK=

Animal Breeding & Genomics Centre

Animal Breeding & Genomics Centre

- Different populations
- Different nutrition and management(-groups)
- Different recording strategies

LIVESTOCK RESEARCH WAGENINGEN UR



Introduction

Objective:

Test the accuracy of DGV from data pooled across 4 research herds (UK, SW, NL, IE), with the progeny test EBV in the UK, IE and NL.

Steps:

- Combine genotypes & phenotypes across herds
- Estimate SNP key and DGVs
- Correlate with progeny breeding values for national traits





Methods	Accuracy Robust	Milk DGV		
 Random regression model with random animal effect across DIM, to adjust for fixed effects and predict full lactation curves for each animal 	animal cts and nal		Correlation with RobustMilk DG (194 bulls)	
	Progeny EBV	NL	IE	UK
2) BSSVS (Calus et al. 2008) model: Gibbs sampling	Milk (kg)	0.51	0.42	0.57
50 000 cycles 5 chains	Fat (kg)	0.57	0.51	0.58
	Protein (kg)	0.46	0.40	0.56
a) DGV/: polygenic plus SNP effects average of 5	Fat (%)	0.78		0.76
chains, i.e. no parent average in there.	Protein (%)	0.71		0.70
4) Correlation DGV and EBV ≈ accuracy DGV				
UVESTOCK RESEARCH WASCHINGENED	LIVESTOCK RESEARCH WAGENINGEN DR	ROB	USTMILK=	Animal Breeding & Genomics Centre

	Correlation with RobustMilk DGV			
	LW (kg)	BCS (1-5)	DMI (kg/d)	
Progeny EBV NL				
Angularity	-0.30	-0.30		
Body condition score	0.40	<u>0.36</u>	0.17	
Chest width	0.41	0.28	0.29	
Dairy strength	0.37	0.26	0.20	

Expected versus observed accuracy



	Accura	Accuracy RobustMilk DGV (44 bulls)		
	NL	IE	UK	NL
Milk(kg)	0.36	0.24	0.48	0.51
Fat(kg)	0.38	0.51	0.31	0.57
Protein(kg)	0.24	0.23	0.35	0.46
Fat(%)	0.79		0.79	0.78
Protein (%)	0.73		0.70	0.71

Conclusion					
 Relatively high accuracy even with Differences in recording and populations Most bulls less than 5 daughters No blending with accurate sire/mgs EBV 					
Lower when no daughters in reference					
Sharing data from research herds might make selection for new traits possible (and effective)!					
	Animal Breeding & Genomics Centre				

Acknowledgements

- The RobustMilk team
- The RobustMilk project is financially supported by the European Commission under the Seventh Research Framework Programme, Grant Agreement KBBE-211708. This publication represents the views of the authors, not the European Commission, and the Commission is not liable for any use that may be made of the information.

