67th Annual meeting of the European Association for Animal Production Belfast, 2016

Towards preventive health management in native dual-purpose cattle via novel breeding strategies

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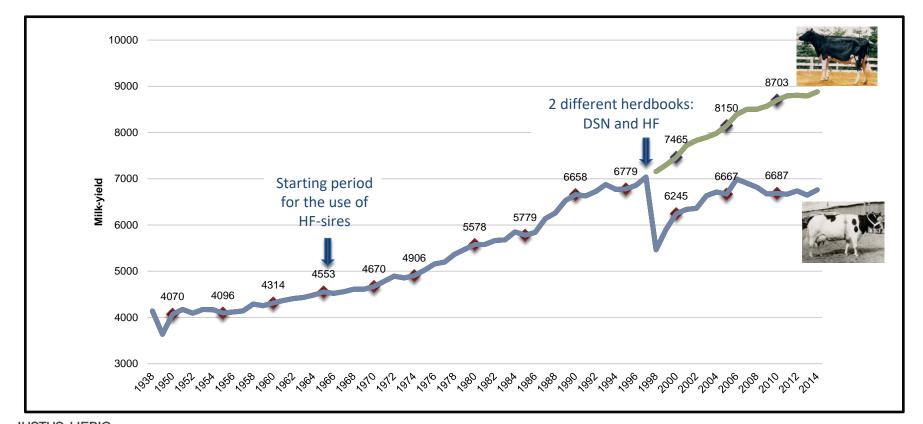
Breeding history

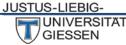


"The development from the dual-purpose DSN breed towards the modern HF cow is the most impressive breeding event in the past 50 years" Dr. P. O. Grothe, former president of the German Holstein Association, 1994



Breed comparison for "conventional" traits





Breed comparison for novel functional traits

Hypothesis:

Dual-purpose cows are more robust than modern dairy cattle breeds, and they are more suitable for <u>harsh organic grassland systems</u> regarding to <u>constitution</u>, <u>physiology</u>, <u>health</u>, <u>milk quality</u> and <u>greenhouse gas emissions</u>



DSN cows in grasland systems from the EU-project: 2-Org-Cows



Simmental cows in grasland systems from the EU-project: 2-Org-Cows

Verification of the hypothesis implies recording of novel functional traits!!!



Installation of Sensor-technology (Longitudinal data for health, rumination, movements,....)

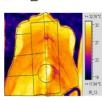


Measurements for methane emissions on an individual basis



Milk fatty acid profiles based on spectral data



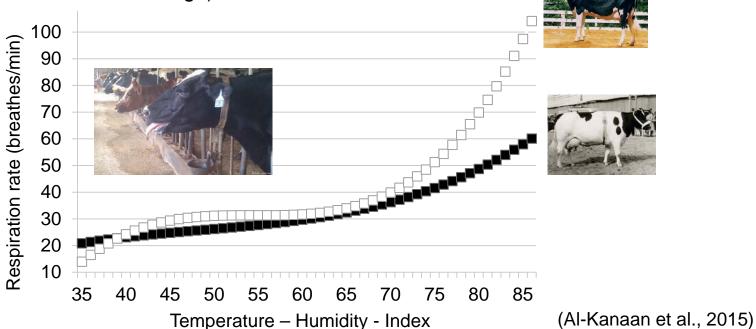


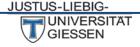
Heat stress indicators: Infrared pictures, respiration rate, body temperature, pulse,...

Breed comparison: Least-Square-Means for respiration rate

83 HF und 155 DSN cows

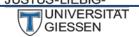
 Data recording from 2012 - 2014 in 2 herds (cross-classified research design)





Second objective: Optimization of within-breed breeding strategies

.....a challenge for endangered breeds because of:



The conventional progeny testing (PT)

Beurteilungskriterium	Optimierungskriterium Zuchtfortschritt Populationsgröße			Züchtungsgewinn			
	100 000	400 000	1 600 000	100 000	400 000	1 60	
Zuchtfortschritt je Kuh			X				
und Jahr in DM	11,15	12,77	14,06	10,62	11,97	13,	
Züchtungsgewinn je Kuh	,						
in DM	31,73	42,35	47,51	37,18	47,75	,54,	
Zuchtaufwand je Kuh							
in DM bezogen auf							
die Gesamtpopulation	26,10	00 00	00 04	10 60	1916	12	
Pay-Off-Periode in Jahren	17,6	Fou	vson a	nd Ni	shal 1	QQ	
Interne Verzinsung in %	9,4	160	vson a	na ivie	SDEI, 1	.70	
Prozentanteil der Milch-							
leistung am Zuchtfortschritt	83,3	80,0	81,7	74,5	76,1	78,	

Tabelle 3. Zuchtfortschritt in Abhängigkeit von dem Testanteil und der Anzahl Bullenv ter für 3 Populationsgrößen

Genetic gain depending on the proportion of cows inseminated with young bulls, t) number of bull sires and the population size

Populations- größe (aktives		Be (aktives anteil		Anzahl Bullenväter je Jahr						
Zuchtma	terial)	(%)	2	4	6	8	10			
100 000 (25 000)		20 30	9,94 10,23	9,46 9,69	8,85 9,06	8,26 8,46	7,71 7,90			
٠.		40 50 60	10,38 10,47 10,51	9,82 9,89 9,93	9,18 9,26	8,59 8,68	8,03 8,15			
		70 80	10,50 10,46	9,94 9,94	9,32 9,36 9,39	8,76 8,83 8,89	8,25 8,34 8,43			
400 000 (100 000)		20 30	11,99 12,32	11,85 12,16	11,64 11,88	11,29 11,52	10,98 11,19			
		Fei	wson and	Niebe	l, 1986)	11,60 11,62 11,60	11,28 11,30			
		80	12,41	12,25 12,14	11,88 11,79	11,55	11,29 11,25 11,19			
(400 000)		20 30 40	13,22 13,60 13,80	13,17 13,54	13,02 13,38	12,87 13,22	12,74 13,08			
		50 60	13,91 13,94	13,74 13,84 13,87	13,56 13,66 13,69	13,40 13,49 13,51	13,25 13,34 13,36			
		70 80	13,92 13,84	13,85 13,77	13,66 13,58	13,48 13,34	13,30			

Tabelle 2. Zuchtfortschritt und Züchtungsgewinn in Abhängigkeit von der Populationsgröße und dem Anteil des aktiven Zuchtmaterials

Genetic gain and profit depending on population size and the proportion of the active population

	Anteil des aktiven Zuchtmaterials (%)								
	10	15	20	25	30	35	40		
	Zuchtfo	rtschritt j	e Kuh/Jahr (I	OMO					
100 000 200 000 400 000 800 000 1 600 000 3 200 000	8,54 9,77 10,83 11,75 12,42 12,95	9,10 10,41 11,53 12,39 13,03 13,57 ngsgewir	9,45 10,80 11,95 12,72 12,72 (Niebel	9,69 11,08 12,16 12,91	9,88 11,28 12,30 13,04	10,03 11,44 12,39 13,12	10,15 11,55 12,45 13,17 13,79 14,34		
100 000 200 000 400 000 800 000 1 600 000 3 200 000	31,10 37,57 42,56 46,53 48,94 50,25	32,17 39,05 44,39 48,03 50,20 51,44	32,45 39,60 45,08 48,20 50,27 51,49	32,35 39,69 44,85 47,81 49,83 51,04	32,02 39,53 44,26 47,12 49,11 50,32	31,55 39,19 43,46 46,25 48,23 49,45	30,98 38,58 42,53 45,28 47,25 48,48		

Tabelle 4. Züchtungsgewinn in Abhängigkeit von dem Testanteil und der Anzahl Bullenväter für 3 Populationsgrößen

Profit depending on the proportion of cows inseminated with young bulls, the number of bull sires and the population size

Populations- größe (aktives	Test- anteil	Anzah					
Zuchtmaterial)	(%)	2	4	6	8	10	12
100 000	20	32,62	31,15	28.28	00.00		_
(25 000)	30	34,04	32,35		25,33	22,40	19,45
	40	34,85	33,06	29,41	26,45	23,53	20,60
~	50	35,29	33,51	30,17	27,38	24,45	21,62
	60	3E 40		30,72	27,96	25,27	22,60 -
,	70		33,76	31,14	28,55	26,04	23,56
	80	35,36	33,84	31,44	29,06	26,77	24,51
-	80	34,98	33,76	31,63	29,50	27,46	25,46
400 000 (100 000)	20	42,59	43,20	42,86	41,44	40,04	38,72
(200 000)				44 33		41,20	39,87
	Nie	hela	ad Ea		1978)	41,75	40,43
		oci ui	14161	wson.	19/81	41,95	
		10,00	*****		0)	41,94	40,69
	70	45,16	45,36	44,19	42,93	41.73	40,74
	80	44,31	44,60	43,58	42,44	41,73	40,62
600 000	20	10.10			22,71	21,00	40,35
(400 000)		46,46	48,90	49,33	49,33	49,20	49.03
-00 000)	30	48,54	51,04	51,45	51,41	51,25	
	40	49,74	52,31	52,71	52,65	52,46	51,05
	50	50,42	53,06	53,46	53,38	53,17	52,24
	60	50,69	53,41	53,80	53,71		52,93
	70	50,56	53,37	53,76	53,65	53,48	53,21
	80	49,99	52,89	53,28	53,16	53,40	53,11
			,	00,20	00,10	52,89	52,58

Genetic evaluations: An across-Europe approach

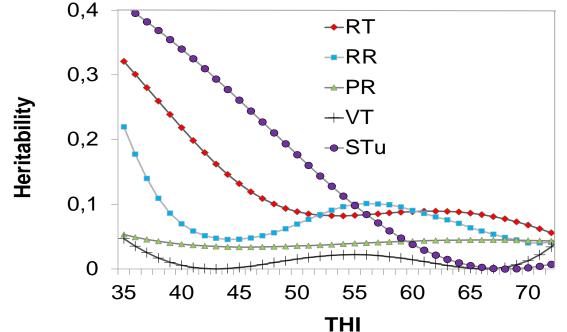
For example for the DSN-breed: Inclusion of herds from Poland, The Netherlands, and Germany



Genetic evalutions: Adaptation and evaluation of alternative methods

A: Genetic parameters in dependency of continuous environmental descriptors, e.g. temperature, humidity, grassland characteristics, etc.

→ Application of random regression models with environmental dependent covariates



(Al-Kanaan et al., 2015)

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Genetic evalutions: Adaptation and evaluation of alternative methods

- B: "borderless clustering" approach
- → Allocation of herds across country borders according to production system characteristics

Based on the idea by Weigel et al. (1999)
Utilisation of production systems for international genetic evaluations instead of country borders!

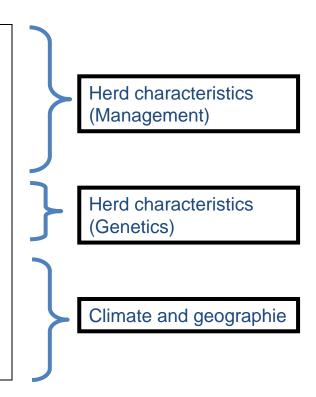
Weigel et al. (2001):

- ❖ 131.9 Mio. test-day data from 16.4 Mio. Holstein cows located in 17 countries
- grouping herds accross country borders into clusters (7 clusters were defined)



Criteria for herd clustering (Weigel et al., 2001)

- 1. Herd size
- 2. Average calving interval
- 3. Milking frquency
- 4. Age at first calving
- 5. 305-d milk yield
- 6. Most frequent month of calving
- 7. Genetic value of sires
- 8. % Holstein genes of sires
- 9. Degree of latitude
- 10. Sea level
- 11. Average temperature in July
- 12. Average rainfall in July
- 13. Percentage of pasture

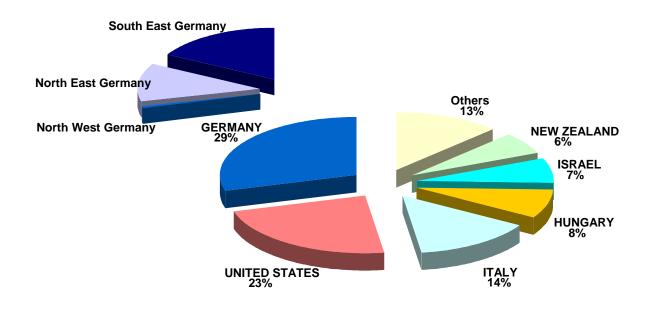


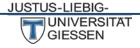
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Cluster 1

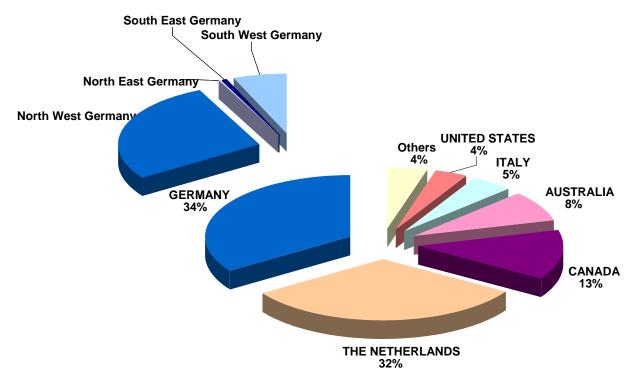
"large herds with high intra-herd SD, high production level"





Cluster 2

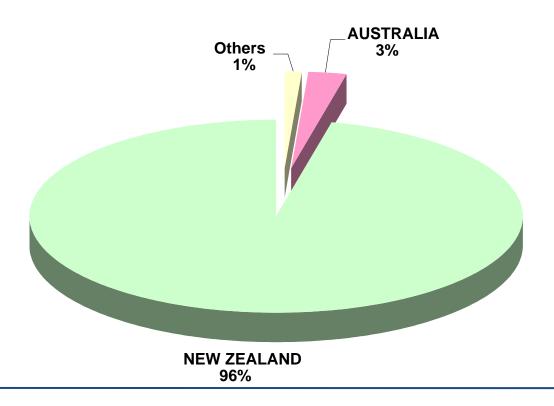
"small herds with low intra-herd SD, high production level"





Cluster 5

"cluster for pasture based herds"





What are the clusters for dual-purpose cattle?

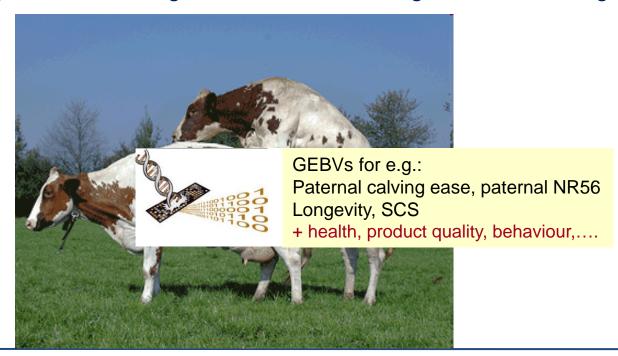
<u>Aim</u>: Identification of relevant environmental and herd descriptors for dual-purpose cattle accross Europe, and clustering herds accordingly

First attempt: See presentation by Maria Jaeger in this session

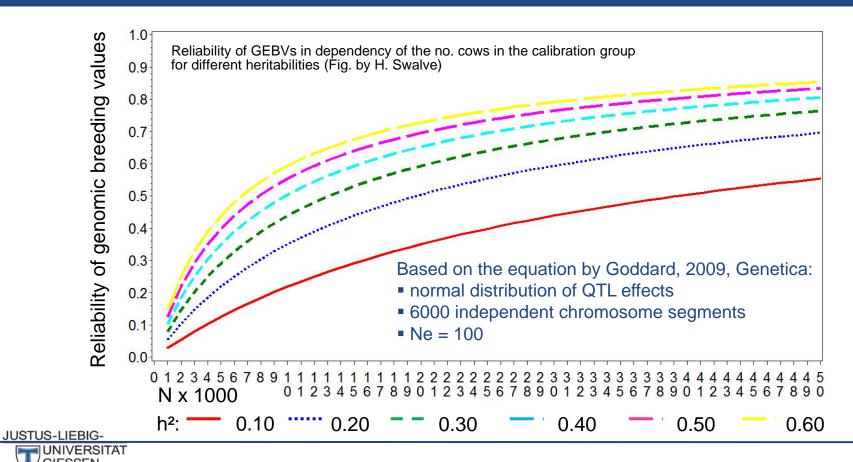


Genetic improvements of novel health traits in dual-purpose cattle

- Impossible to setting up the infrastructure and logistics for a PT-program
- Alternative: Implementation of genomic selection using cow calibration groups



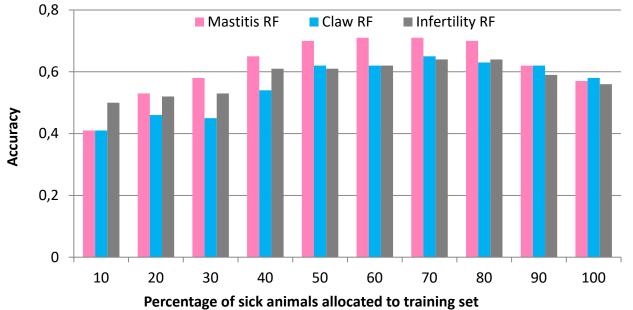
Cow calibration groups: How many cows should be included?



The German cow calibration group project for health traits in HF cattle

- 20'000 genotyped Holstein cows and their imputed dams
- Health trait recording according to a central diagnosis key (956 single disease traits)

Accuracy of genomic predictions for laminitis, clinical mastitis and infertility (Naderi et al., 2016)



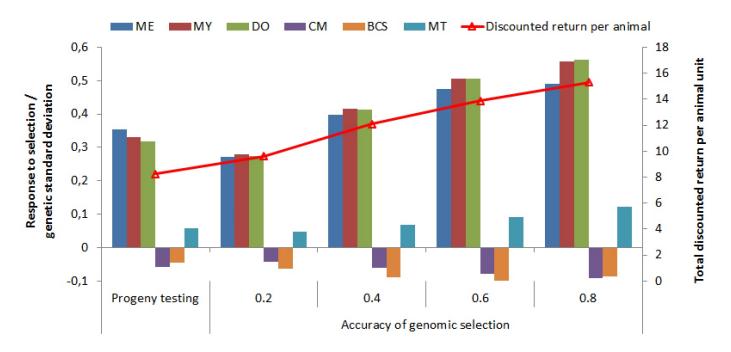


....but we do not have 20'000 DSN cows with pheno- and genotypes

- Solution A: Using SNP-effects from HF-calibration groups for predictions in DSN
 - usually this does not work
- Solution B: Mixing breeds in reference groups
 - → acceptable results for a similar no. of cows from both breeds
- Due to shared founder animals, genetic relationships between DSN- and HF-populations do exist (< 1%).
 - → further increase of reliabilities of genomic breeding values for DSN cows is expected
- All project partners intend to genotype a fraction of dual-purpose cattle
 - cow calibration group across country borders!
- Studie von Van Eijndhoven et al. (2015), J. Dairy Sci. 98, 6510-6521
 High correlations between SNP effects contributing to milk quality traits when using DSN- and HF-cows

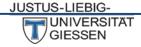


Definition of breeding goals / evaluation of breeding strategies



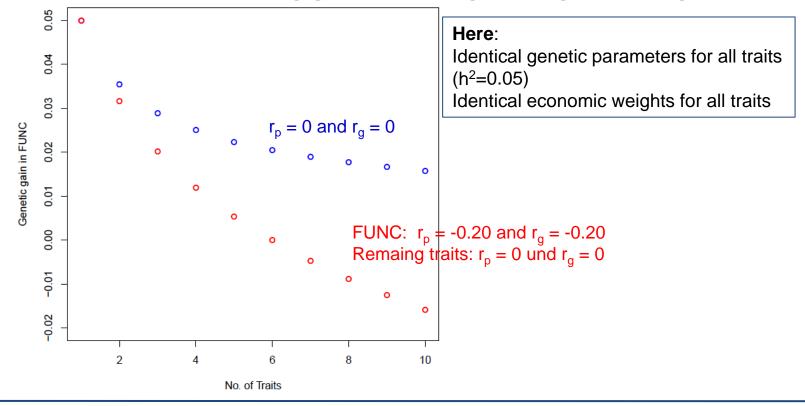
Proeny testing and genomic breeding program with different accuracy

(Yin et al., 2015)



The breeding challenge

An increase of traits in the breeding goal reduces genetic gain in single traits!!



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Conclusions: Steps for preventive health management via breeding in dual purpose cattle kept in "harsh environments" (grassland systems)

- 1. Recording of novel functional traits (traits of interest and indicator traits)
- 2. Developing and evaluating alternative models for genetic evaluations (Consideration of GxE-interactions, e.g. RRM or "borderless clustering"
- 3. Setting up cow calibration groups on a European scale
- 4. Defining overall breeding goals and evaluating alternative breeding strategies



Thanks in advance to all project partners for their scientific contributions in order to realize those challenging breeding tasks!

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