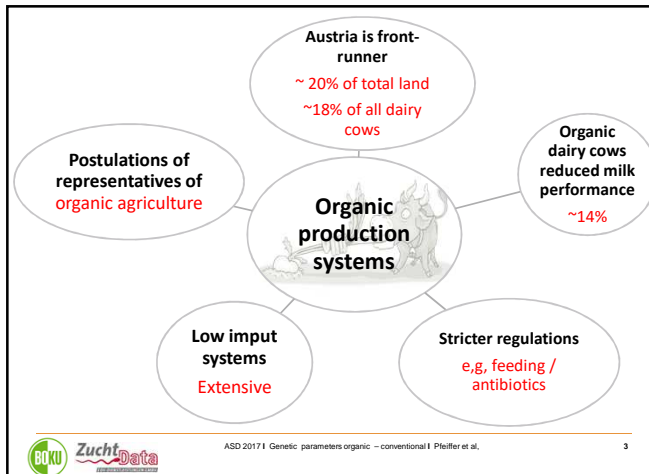
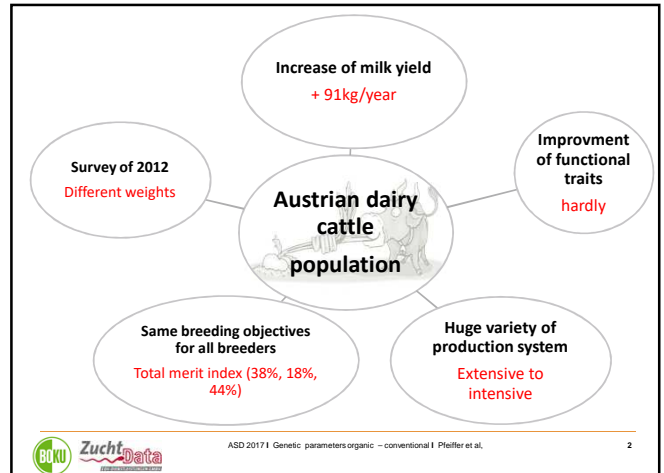


# Genetic parameters of Austrian Fleckvieh cattle in organic and conventional production systems with different levels of management

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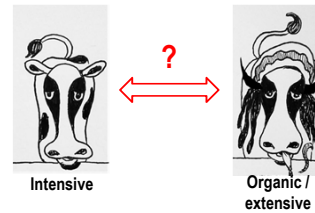
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## Assumptions

- Organic and extensive farming strategies follow different breeding goals



→ Do we select the best animals for the different production systems in Austria?

## Objectives (I)

- Investigation of G x E interaction for different production systems (Pfeiffer et al., 2016 Livestock Science)
- Estimation of genetic parameters ( $h^2$  and  $r_a$ ) of different production systems
  - Milk kg
  - Fat kg
  - Protein kg
  - Persistency
  - Somatic cell score (SCS)
  - Functional longevity (LONG)
  - Milking speed (MS)
  - Udder health index (UHI)
  - Fertility index (FI)



## Objectives (II)

- Conduct an approximate multitrait 2-step procedure applied to yield deviations and de-regressed breeding values
- According to results recommendations for specific breeding programmes can be given



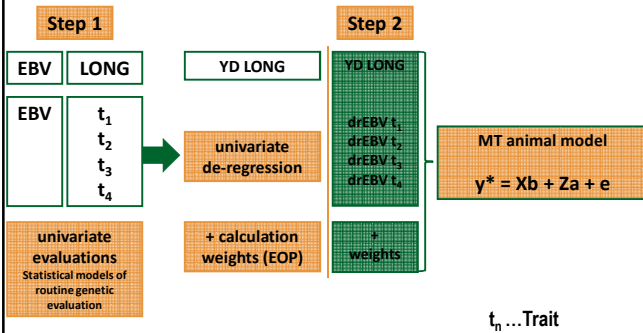
## Materials & methods

- Data basis
  - Whole Austrian Fleckvieh cow population
  - Maximum 12.5% non-Fleckvieh-gene proportion
  - All 9 traits recorded
  - Born between 1999 and 2011
  - Known parents
  - Herd size  $\geq 5$  lactating cows
- Organic (Organic) - conventional low (ConL) – conventional high (ConH)
- Grouping according to the herd-year-effect (HYE) for milk yield
  - $\pm 2$  SD from the HYE mean
  - Indicator for farm management

## Summary of the production systems

	Organic	ConL	ConH
Animals (n)	46,015	30,840	53,882
Farms (n)	2,657	1,948	1,486
Range HYE	3,274 – 7,287	1,948 – 4,800	6,220 – 8,350
Mean HYE	5,076	4,452	6,558
SD HYE	574	305	312
Pedigree (n)	162,560	118,010	159,943

## Approximate multitrait 2-step procedure



## Models

- Univariate or bivariate animal model

$$y^* = Xb + Za + e$$

- ASReml

## Results and Discussion – Heritabilities

	Organic	ConL	ConH
Milk kg	0.63	0.59	0.65
Fat kg	0.50	0.45	0.55
Protein kg	0.57	0.51	0.57
Persistency	0.56	0.52	0.62
SCS	0.23	0.22	0.29
LONG	0.07	0.09	0.04
MS	0.27	0.21	0.33
UHI	0.12	0.15	0.21
FI	0.06	0.04	0.05

SE  $\leq 0,02$

## Results and Discussion – Genetic correlation MILK kg

	Organic	ConL	ConH
Fat kg	0.66	0.70	0.66
Protein kg	0.84	0.85	0.86
Persistency	-0.04*	-0.12	0.01*
SCS	-0.01*	0.09	0.01*
LONG	-0.23	-0.29	-0.23
MS	0.36	0.45	0.34
UHI	-0.01*	0.05*	0.01*
FI	-0.15	-0.32	-0.14

\*not significantly different from zero

## Conclusions

- Applied procedure was **feasible**
- **Similar genetic heritabilities** and **genetic correlations** between traits
- **Similar** production environments and farm managements
- **Slight G x E** for **several traits but close to unity** (Pfeiffer et al., 2016)
- **Independent breeding objectives** are currently **not needed**
- More research required for **indicators of ConL farms**



ASD 2017 | Genetic parameters organic – conventional | Pfeiffer et al.

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