

# Prediction of Breeding Values and Aggregate Genotype

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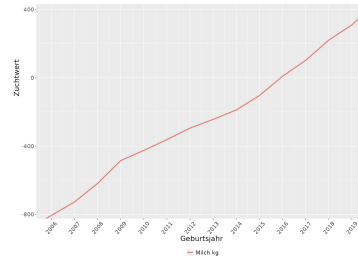
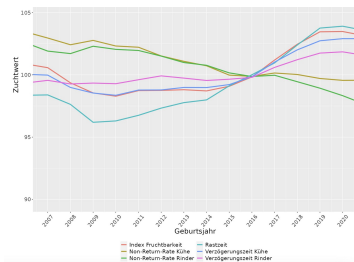
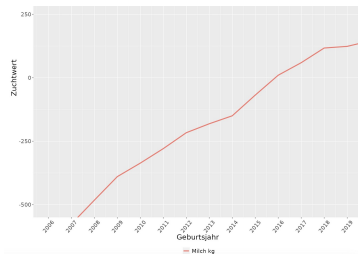
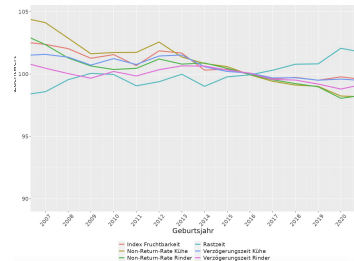
2024-05-15

# Summary for One Trait

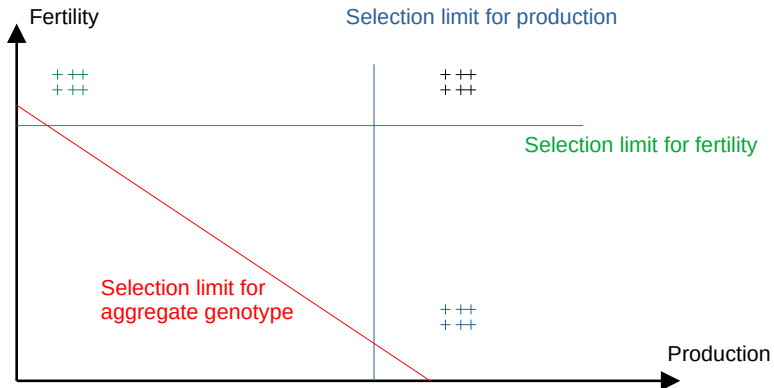
- ▶ Predicted breeding values with or without genomic information
- ▶ Animals can be ranked according to predicted breeding values
- ▶ Problems:
  - ▶ Not only one trait should be improved
  - ▶ Selection for one trait changes also other traits via correlated selection response

→ see genetic trends at: <https://1-htz.quagzws.com/shiny/users/zws/genTrendHO/DE/2024/>

# Example Fertility and Production



# Multi-Trait Selection



# Types of Multi-Trait Selection

- ▶ Tandem selection
  - ▶ select for one trait at the time
  - ▶ after goal has been reached change to different trait
- ▶ Independent selection limits
  - ▶ select only animals which fulfill criteria in all traits
- ▶ Selection according to aggregate genotype
  - ▶ combine traits into aggregate genotype  $H$
  - ▶ define  $H$  as weighted sum of true breeding values and economic values
  - ▶ use selection index  $I$  to estimate  $H$

# Aggregate Genotype

Definition in vector notation:  $H = v^T \cdot u$

where

- ▶  $u$ : vector of true breeding values
- ▶  $v$  vector of economic values which are marginal changes in profit for a small change in the population mean of the trait

Estimate  $H$  via index  $I$ , hence  $\hat{H} = I = b^T x$

with

- ▶  $x$ : a vector of information sources
- ▶  $b$ : a vector of unknown weights.

Determine  $b$  such that  $\text{var}(I - H)$  is minimal.

Find  $b \dots$

$\dots$  such that  $\text{var}(I - H)$

$$\begin{aligned}\text{var}(I - H) &= \text{var}(I) - 2 * \text{cov}(I, H) + \text{var}(H) \\ &= \text{var}(b^T x) - 2 * \text{cov}(b^T x, v^T u) + \text{var}(v^T u) \\ &= b^T \text{var}(x) b - 2 * b^T \text{cov}(x, u^T) v + v^T \text{var}(u) v \\ &= b^T P b - 2 * b^T C v + v^T G v\end{aligned}$$

Setting  $\frac{\partial \text{var}(I-H)}{\partial b} = 0$  leads to

$$Pb = Cv$$

Hence

$$b = P^{-1}Cv$$

## Special Case

- ▶ Same traits in  $H$  and in  $I$
- ▶ Use predicted breeding values  $\hat{u}$  from multivariate BLUP animal model as information source  $x$
- ▶ Then it follows

$$b = P^{-1}Cv = \text{var}(\hat{u})^{-1} \cdot \text{cov}(\hat{u}, u^T) \cdot v = v$$



## Method Works . . .

### Studies like

- ▶ Berry, D. P., Kearney, J. F., Twomey, K., & Evans, R. D. (2012). Genetics of reproductive performance in seasonal calving dairy cattle production systems. *Irish Journal of Agricultural and Food Research*, 52(1), 1–16.