



IT-Solutions for
Animal Production



Horse Evaluation Webinar / EquiMais on 4 June 2021:

'Importance of the morpho-functional evaluation in the genetic improvement of horses'

Equine genetic evaluation with special emphasis on linear description

Kathrin F. Stock

IT Solutions for Animal Production (vit), Verden, Germany

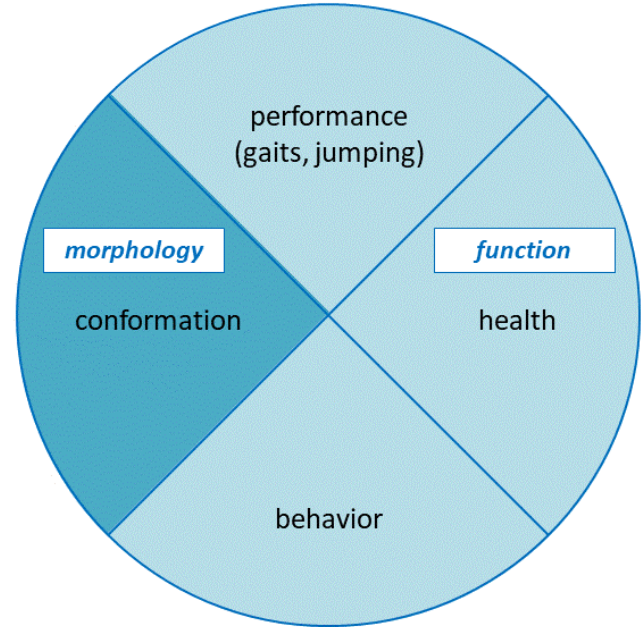
Outline

- ❖ breeding goals and breeding programs of sport horses
 - ◆ data recording
 - ◆ data analyses (incl. genetic evaluation)
- ❖ routine linear profiling and genetic evaluation for linear traits
 - ◆ advances and challenges for horse breeding
 - ◆ data structure, traits and modelling
 - ◆ role of genetic linear profiles
- ❖ implications for breeding applications



Breeding goals of sport horses

- centered on performance under saddle
- different aspects (interactions)
- meeting the needs of the equine sector
 - specialists rather than generalists
 - internationally competitive horses
 - for professional and amateur riders
 - from top sport to leisure riding
- more diverse and more challenging settings than in other livestock species



Breeding programs & data recording

- systematic data recording as basic requirement in animal breeding
- the earlier and the more reliable the data recording, the better the chances of lasting success of breeding programs
- clear target definition as key factor for success (= breeding progress)
 - direct measures of breeding goal traits,
e.g. competition performance on highest level, lifetime performance, placings in dressage competitions, ranking points in show jumping competitions
 - indirect measures (indicator traits)
e.g. quality of gaits, jumping ability and style, functionally important conformational aspects, absence of radiographic findings of presumed clinical relevance



Data analysis & use: general aspects

- **high quality data recording** as important first step to be followed by steps of data processing, data analyses, results presentation and use
- influences on the value of the outcome for breeding applications:
 - evaluated sample of horses
(age, representative of the population)
 - evaluation method
(information value, reliability, objectivity, proximity to the breeding goal)
 - data distribution and modelling
(phenotypic variability, reflected differences between individual horses)
 - transparency and overall acceptance of the system
(trust in the whole system from data recording to selection support tools)



Data analysis & use: specific aspects

- **high quality data recording** as important first step to be followed by steps of data processing, data analyses, results presentation and use
- framework of equine breeding applications:
 - very different types of breeders → challenging job of studbooks (heterogeneous target group with wide range of background, knowledge, ambitions)
 - often emotional rather than rational decision making
 - common skepticism towards more sophisticated analyses (departure from phenotypes) → difficult stand and limited use of breeding values

Genetic evaluations for sport horses I

- based on two main sources of data
 - routine assessments of studbooks
 - registration, studbook entry, performance test
 - traditionally valuating scores, increasingly linear profiles
 - sport data
 - incl. young horse competitions (valuating scores)
 - from regional to national, increasingly international
- PROs and CONs
 - data quality, information value (specificity of traits), preselection bias
 - proximity to the breeding goal



Genetic evaluations for sport horses II

- in Germany two types of genetic evaluations:
 - (1) national genetic evaluation for riding horses (FN-ZWS)**
 - across studbooks
 - sport traits (rank-based and highest level achieved) and young horse traits (valuating scores)
 - dressage and show-jumping
 - breeding values for 12 individual traits + 2 young horse indices
 - positive feedback to the two newly introduced traits (since 2019), HEK Dressage and HEK Jumping, reflecting the 'ultimate breeding goal'
→ major disadvantage: late availability of the definite phenotype

Genetic evaluations for sport horses II

- in Germany two types of genetic evaluations:
 - (1) national genetic evaluation for riding horses (FN-ZWS)
 - (2) supplementary genetic evaluations
 - within studbook
 - conformation and performance (gaits, jumping)
 - different basis: valuating scores (HANN, HOL, TRAK, DSP / BrbA) or linear profiles (OL/OS); mostly young horses $\geq 2 \frac{1}{2}$ years of age (mares, stallions +/-, geldings +/-), sometimes also foals
- different portfolios of breeding values and indices (per studbook)
- optimum conditions to be best fit solution for respective breeders



Horse evaluation: requirements I

- data recording: informative evaluation criteria
 - clearly and unambiguously defined
 - comparable, repeatable (under field conditions)
 - informative (individual horse, population)
 - breeding goal related, early assessable
- preconditions for data use in breeding:
reliable measure or indicator of underlying genetics

objective
precise
specific
decision-relevant



Horse evaluation: requirements

- data recording: informative evaluation criteria
 - clearly and unambiguously defined
 - comparable, repeatable (under field conditions)
 - informative (individual horse, population)
 - breeding goal related, early assessable
- preconditions for data use in breeding:
reliable measure or indicator of underlying genetics

objective
precise
specific
decision-relevant

traditional valuating scoring

- judgement relative to the breeding goal
- subjective scoring scale
'very bad' (1) to 'excellent' (10)

linear description (linear profiling)

- assessment relative to biological extremes
- linear scale independent from optimum expression
(e.g.. 'very short' to 'very long', 'very low' to 'very high')

Horse evaluation: requirements II



*strong in type, much gender expression,
noble head, high-set neck, ...,
trot with much knee action, energetic,
very supple, with active hind limbs, ...*

*overall score of
8.0 or 8.5 or ...?*

*objective
precise
specific
decision-relevant*

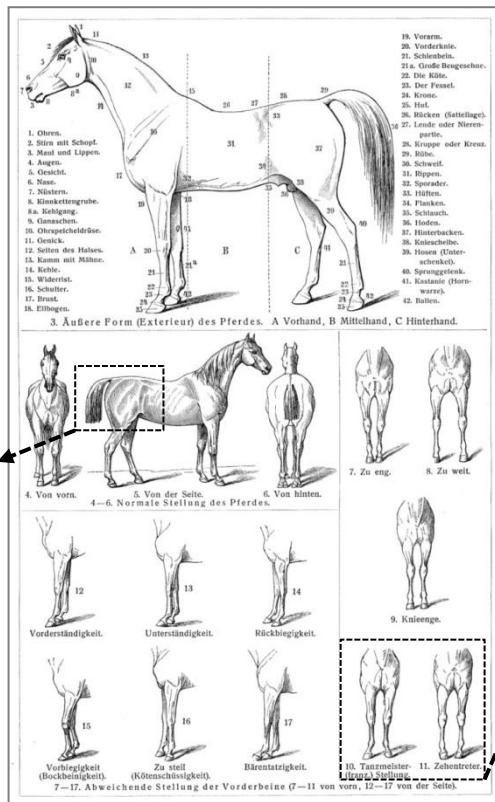
traditional valuating scoring

- judgement relative to the breeding goal
- subjective scoring scale
'very bad' (1) to 'excellent' (10)

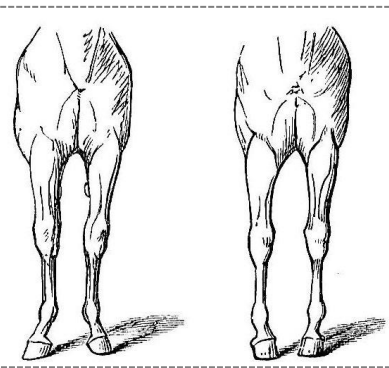
linear description (linear profiling)

- assessment relative to biological extremes
- linear scale independent from optimum expression
(e.g.. 'very short' to 'very long', 'very low' to 'very high')

Horse evaluation: requirements III



**description of specific aspects
as basic principle of horse evaluation**
→ linear description (or linear profiling)
as practice-proven way to do it systematically

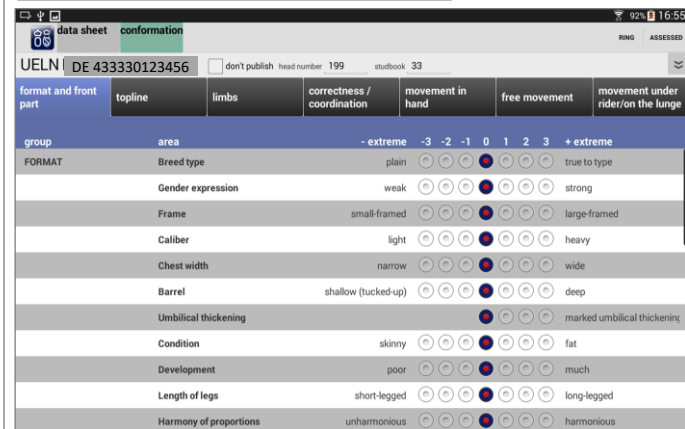
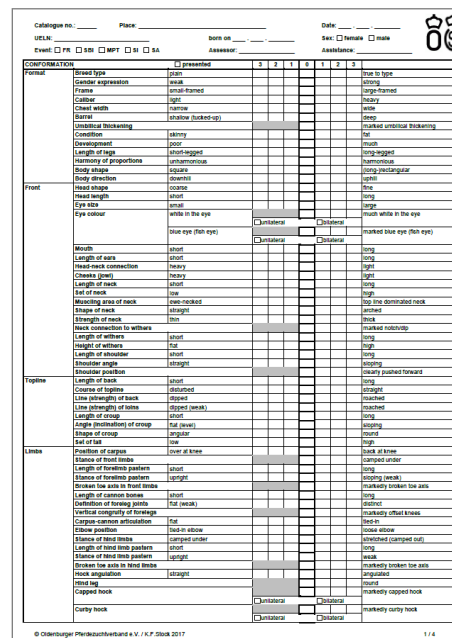


source: Meyers Großes Konversations-Lexikon (Band 15), Leipzig 1908.

Linear description: motivation & start

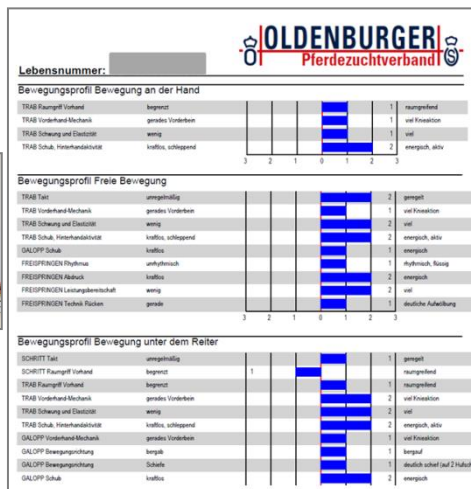
- new role of phenotyping and data-based decision support
 - need of specific information for targeted breeding work
 - advanced methodology requiring high quality data
- **interest in genomic selection / genomic applications in horses**
increasing awareness of the importance of horse evaluation
- obvious limitations of the traditional valuating scoring system vs. long-known advantages of linear recording systems
- **science-to-practice / practice-to-science initiative**
to implement linear profiling in sport horse breeding





- pilot studies related to research projects (extra staff for testing and optimizing the system, first data screening)
- application-oriented project work

→ **sound basis for broad implementation**



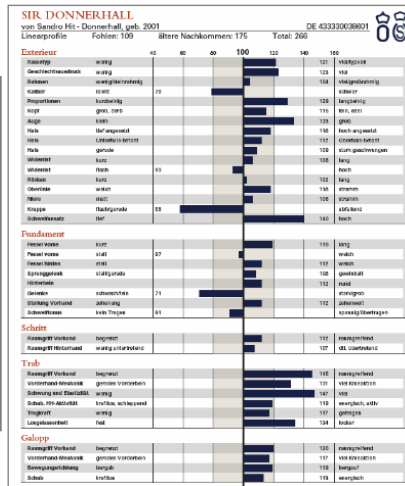
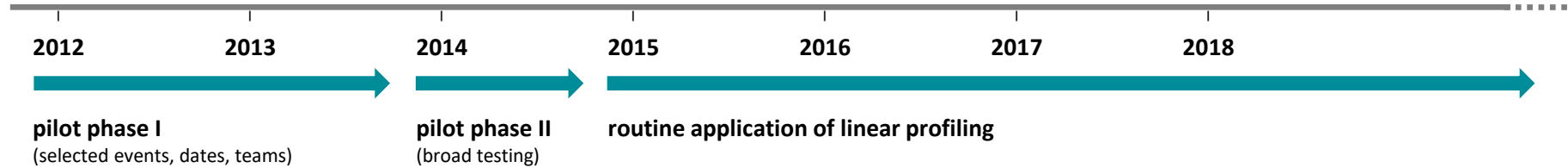
genetic profiles (stallions)

- monitoring of linear data collection and data quality management
 - quality and coverage of the linear scheme (lacking traits?)
 - use of traits and scale, internal consistency of linear data, ...
(unclear definitions? topics for training sessions and technical meetings)
- consolidation and improvement of applications

→ **smooth transition** to new routine applications (acceptance↑)

→ **improved services** for the studbooks and for individual breeders

Linear profiling in Oldenburg (OL, OS) |||



individual linear profiles (phenotypic profiles; foals, mares)

genetic profiles (stallions)

(B) routine and extension

- monitoring of linear data collection and data quality management
 - quality and coverage of the linear scheme (lacking traits?)
 - use of traits and scale, internal consistency of linear data, ...
(unclear definitions? topics for training sessions and technical meetings)
 - consolidation and improvement of applications
- **smooth transition** to new routine applications (acceptance↑)
- **improved services** for the studbooks and for individual breeders

Determinants of the development

- **close collaboration of science and practice**
as key to fast progress and successful implementation
 - smooth transition from R&D to routine
 - transparency and exchange, flexibility of applications
→ acceptance and use

- **early use of linear data** as important driver of innovation
 - continued engagement for optimization of the whole system
(data quality management)
 - high motivation of maximize information output
(internal use → publication of consolidated information)

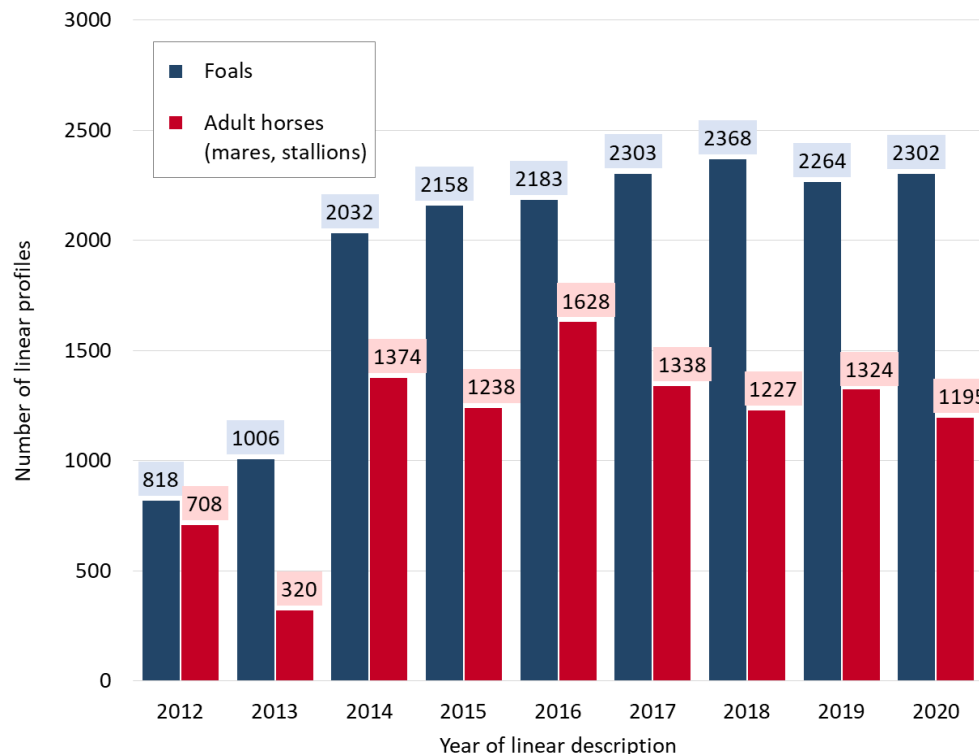


Linear data basis 2012 - 2020

- Oldenburg linear scheme (OL, OS)
 - conformation and performance (gaits, jumping, behavior)
 - 7-point numeric linear scale (-3 to +3)
defect traits / remarks: reduced scale (0 to +3)
 - foals, mares, stallions
(registration, studbook inspection, mare performance test, preselection for licensing, selected young horse competitions)
- extension of linear data collection (routine)
 - most/all events since 2015 in Germany and abroad
 - kernel team of experienced assessors
- **in total N=27,786 linear profiles of 26,069 horses**



Distribution of linear data




- 27,786 linear profiles
(foals : adult horse $\approx 2 : 1$)
- 26,069 horses
incl. 1,608 horses linearly described as foal and adult horse
- 2868 sires
incl. 800 sires with linearly described foals and adult horses (progeny in both age groups)
- optimum use of linear data across age groups



Data collection → Genetic analyses

- mobile data collection (tablet PC) making it possible to:
 - efficiently collect detailed linear data (active input of deviations only)
 - routinely work with refined trait definitions and comprehensive linear schema (basis of highly valued individual assessment reports, considerable potential for research)
 - directly use maximal information from routine assessments

➤ **standardized detailed information on: conformation, gaits, jumping, behavior**

 Linear Profiling

format and front part	topline	limbs	correctness / coordination	movement in hand	free movement	movement under rider/on the lunge
group	area		- extreme	-3 -2 -1 0 1 2 3	+ extreme	
TOPLINE	Length of back		short	⊖ ⊖ ⊖ ⊖ ⊕ ⊕ ⊕ ⊕	long	
	Course of topline		disturbed	⊖ ⊖ ⊖ ⊖ ⊕ ⊕ ⊕ ⊕	straight	
	Line (strength) of back		dipped	⊖ ⊖ ⊖ ⊖ ⊕ ⊕ ⊕ ⊕	roached	
	Line (strength) of loins		dipped (weak)	⊖ ⊖ ⊖ ⊖ ⊕ ⊕ ⊕ ⊕	roached	
	Length of croup		short	⊖ ⊖ ⊖ ⊖ ⊕ ⊕ ⊕ ⊕	long	
	Angle (inclination) of croup		flat (level)	⊖ ⊖ ⊖ ⊖ ⊕ ⊕ ⊕ ⊕	sloping	
	Shape of croup		angular	⊖ ⊖ ⊖ ⊖ ⊕ ⊕ ⊕ ⊕	round	
	Set of tail		low	⊖ ⊖ ⊖ ⊖ ⊕ ⊕ ⊕ ⊕	high	

format and front part	topline	limbs	correctness / coordination	movement in hand	free movement	movement under rider/on the lunge
group	area		- extreme	-3 -2 -1 0 1 2 3	+ extreme	
CORRECTNESS/	Toe stance of forelegs		toe-in	⊖ ⊖ ⊖ ⊖ ⊕ ⊕ ⊕ ⊕	toe-out	
SPECIAL REMARKS	Standing position of front li		base-narrow	⊖ ⊖ ⊖ ⊖ ⊕ ⊕ ⊕ ⊕	base-wide	
			unilateral	⊖ ⊖ ⊖ ⊖ ⊕ ⊕ ⊕ ⊕	bilateral	
	Position of carpus - frontal		wide at knees (bow-legged)	⊖ ⊖ ⊖ ⊖ ⊕ ⊕ ⊕ ⊕	narrow at knees (knock-knee)	
	Toe stance of hind legs		toe-in	⊖ ⊖ ⊖ ⊖ ⊕ ⊕ ⊕ ⊕	toe-out	
			unilateral	⊖ ⊖ ⊖ ⊖ ⊕ ⊕ ⊕ ⊕	bilateral	

Genetic analyses

- trait definition
within age group and trait category

- selected linear traits

(standard deviation, variance, kurtosis)

- estimation of genetic parameters
- genetic evaluation
= prediction of breeding values

Trait category	No. of traits	
	recorded	gen. eval.
Conformation	73	23
Walk (H, F, R)	6	2
Trot (H, F, R)	11	6
Canter (F, R)	10	4
Jumping (F)	16	10
Special remarks (H, F, R)	8	1
Behavior (H, F, R)	7	0

H = in hand, F = free, R = under rider

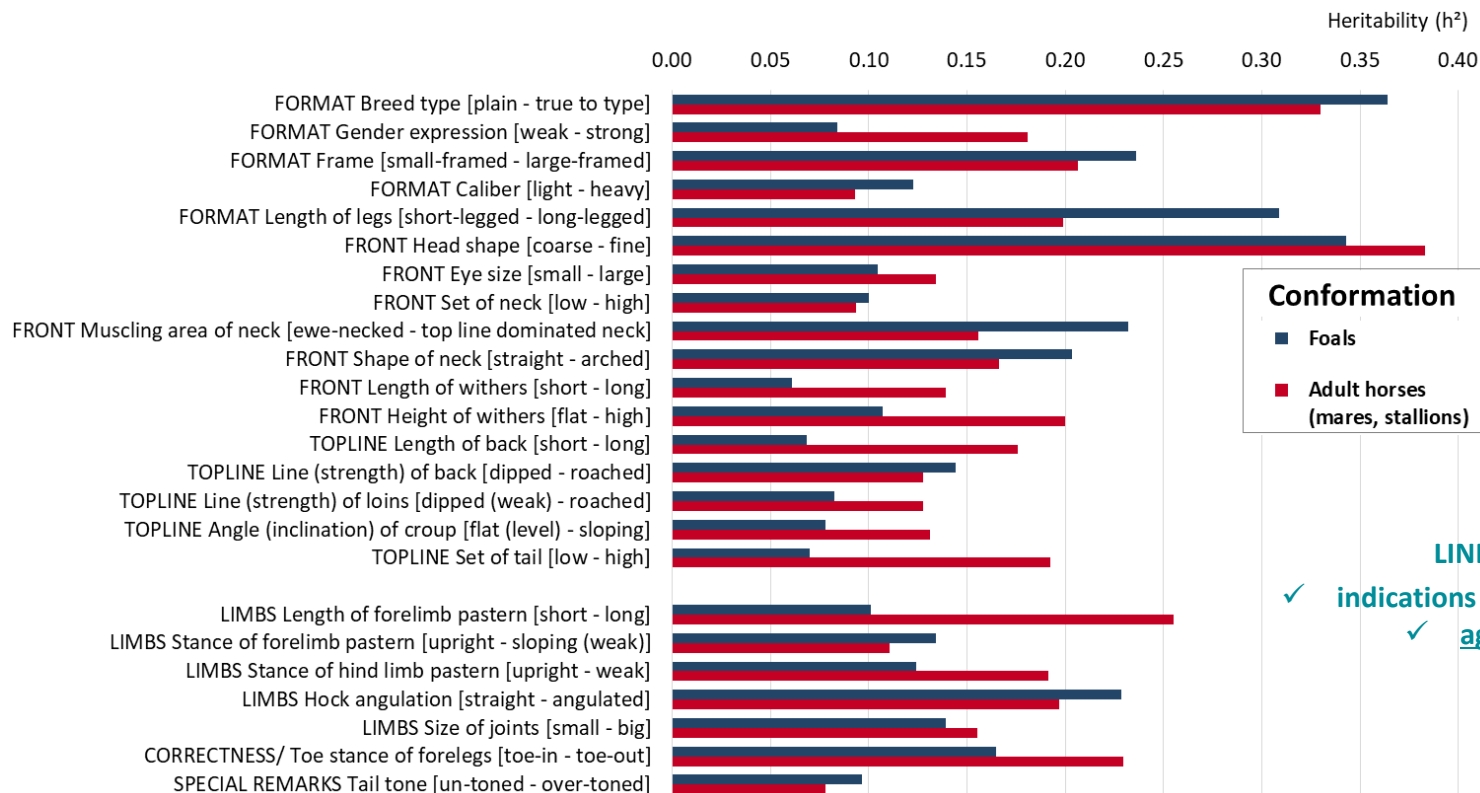
- single- and multiple-trait linear animal models

foals:
$$y_{ijkno} = \mu + SB_i + EVENT-TEAM_j + AGE_M_k + SEX_l + animal_o + e_{ijklop}$$

mares+stallions:
$$y_{ijmnp} = \mu + SB_i + EVENT-TEAM_j + AGE_J_m + PTYPE_n + animal_o + pe_o + e_{ijmnp}$$



Conformation: heritabilities



Conformation

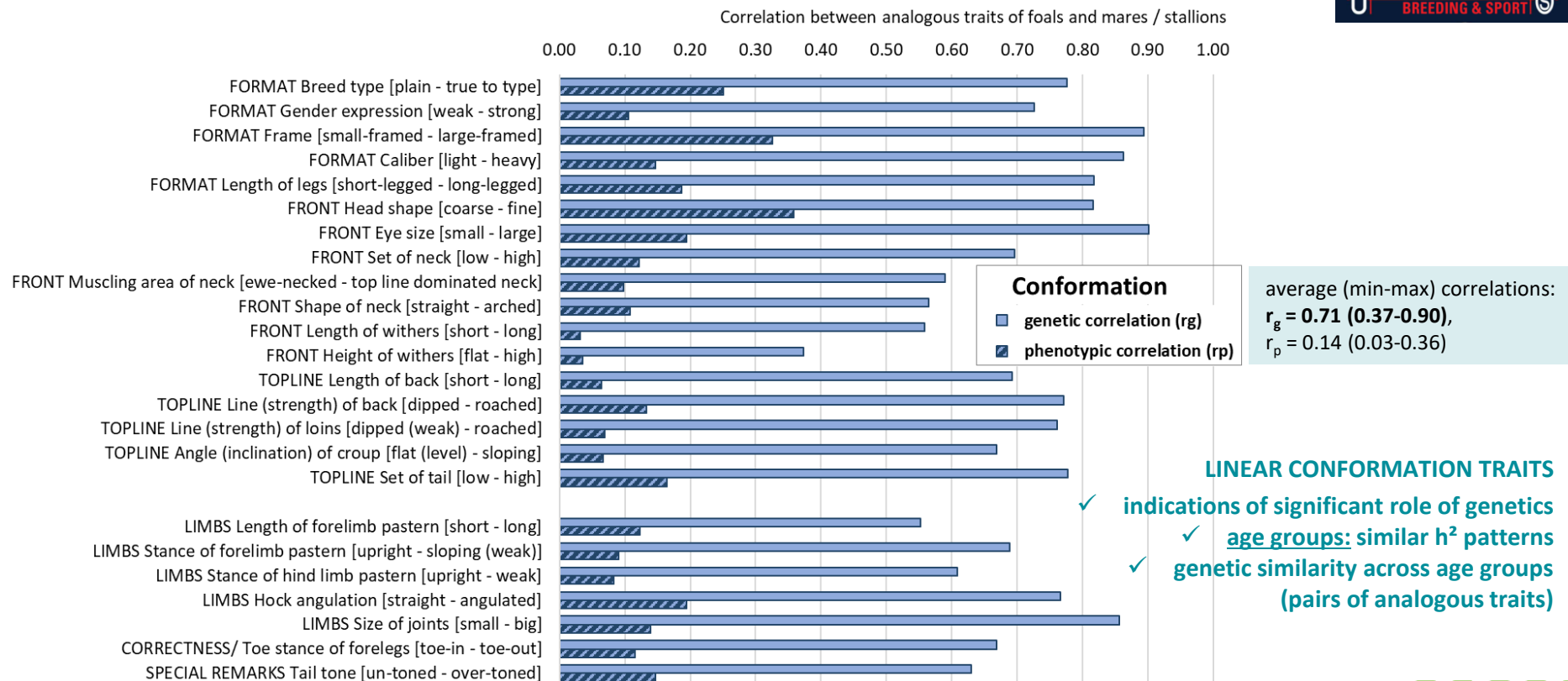
- Foals
- Adult horses (mares, stallions)

average (min-max) h^2 :
 0.15 (0.06-0.36) in foals,
 0.18 (0.08-0.38) in adults

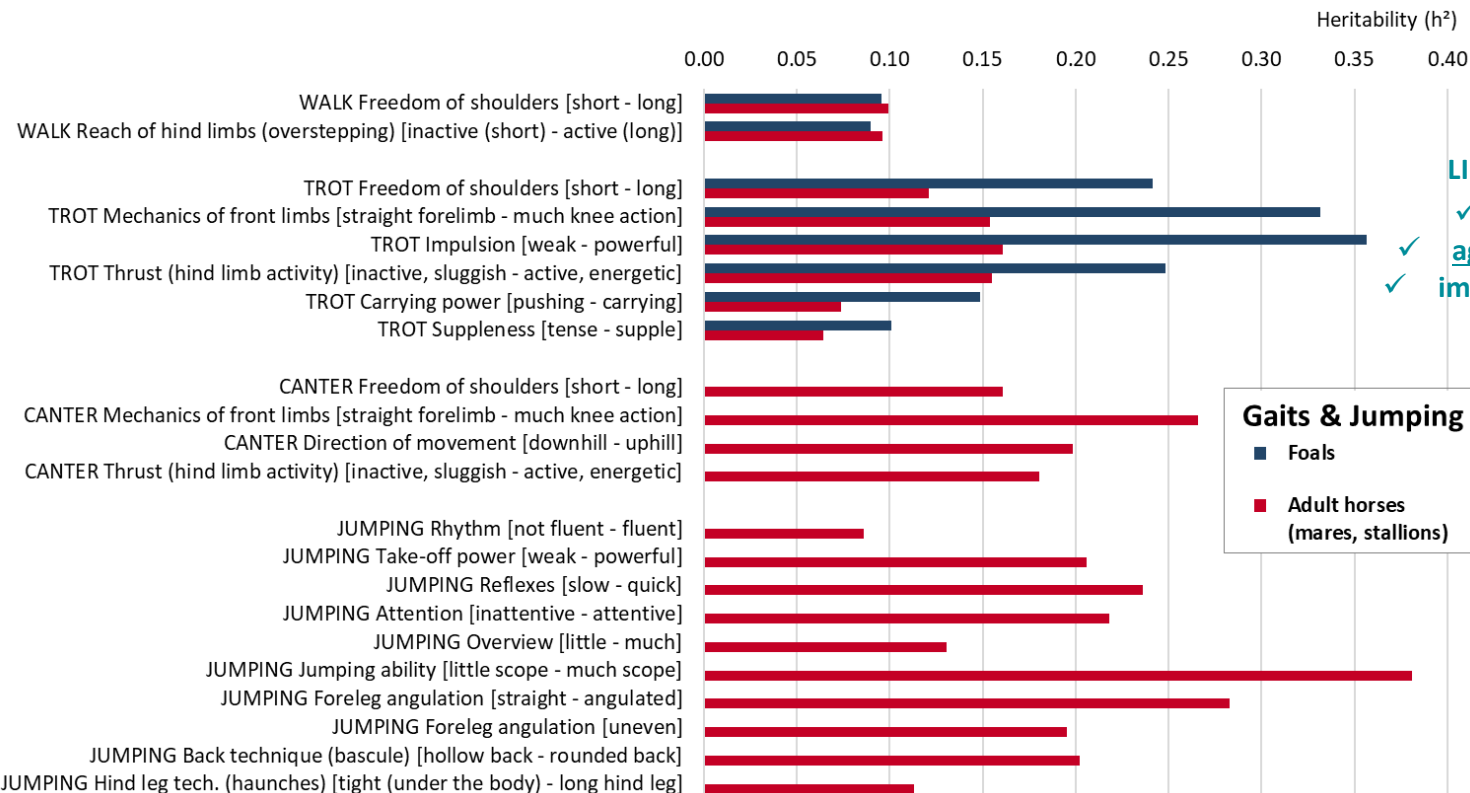
LINEAR CONFORMATION TRAITS

- ✓ indications of significant role of genetics
- ✓ age groups: similar h^2 patterns

Conformation: correlations



Performance: heritability



LINEAR PERFORMANCE TRAITS

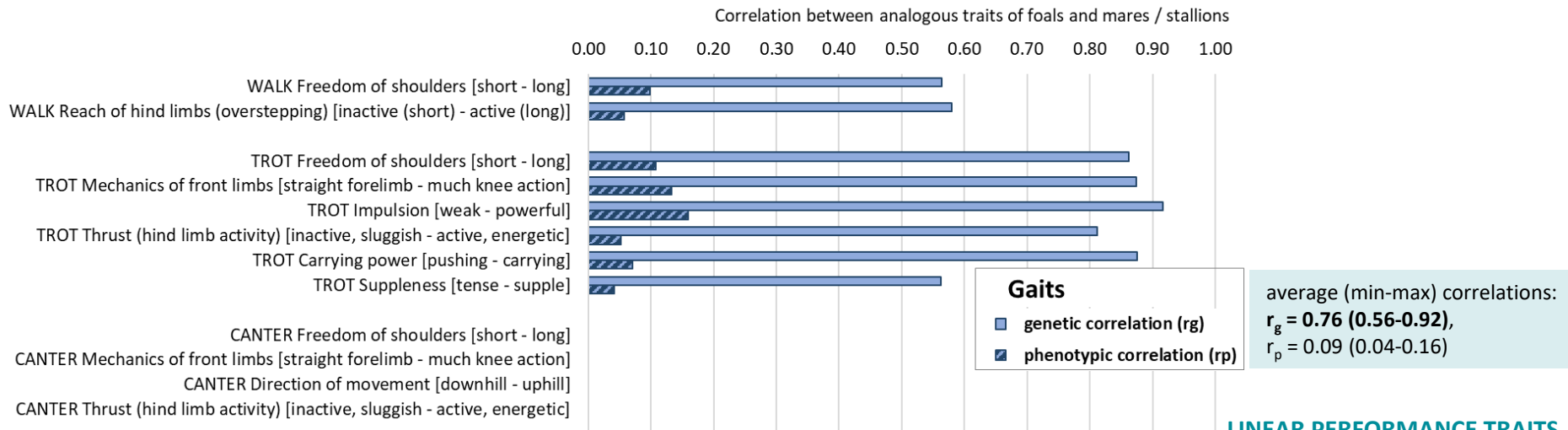
- ✓ important role of genetics
- ✓ age groups: similar h^2 patterns
- ✓ impact of evaluation conditions (information density)

Gaits & Jumping

- Foals
- Adult horses (mares, stallions)

average (min-max) h^2 :
 0.20 (0.09-0.36) in foals,
 0.17 (0.06-0.38) in adults

Performance: correlations



LINEAR PERFORMANCE TRAITS

- ✓ important role of genetics
- ✓ age groups: similar h^2 patterns
- ✓ impact of evaluation conditions (information density)
- ✓ genetic similarity across age groups (pairs of analogous traits)

Data structure & modelling

- results of multivariate estimation of genetic parameters indicating
 - similar patterns of heritabilities (linear aspects with higher/lower h^2)
 - genetic similarity across age groups (pairs of analogous traits)
 - genetic similarity across presentation types (adult horses / different conditions)
- multiple-trait repeatability linear animal model (pairs of traits)
- **high value of linear foal data (broad basis, early prediction)! ✓**



Genetic evaluation: development

- implementation of linear systems for riding horses
 - conformation and performance (gaits, jumping, behavior)
 - foals and/or adult horses (broodmares, stallions, young riding horses)
 - substantial improvement of phenotype data quality: phenotypic linear profiles of individual horses

Linear data basis in the Oldenburg studbooks (OL, OS)

- ✓ since 2012 (end of 2020: 27,786 linear profiles of 20,069 horses)
- ✓ conformation and performance (movement, jumping)
- ✓ same linear scheme and 7-point numeric linear scale (-3 to +3) for all horses, i.e. across age groups

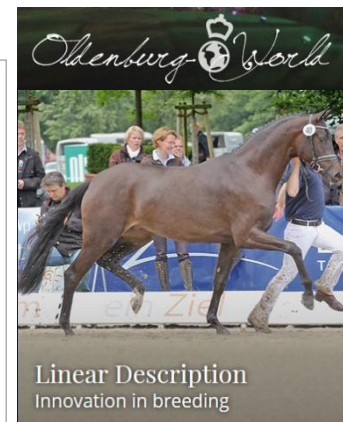


Genetic evaluation: development II

- implementation of linear systems for riding horses
- extension of breeding applications based on linear profiling
 - clearly distinct set of traits (descriptive)
 - potentially powerful tool for breeders

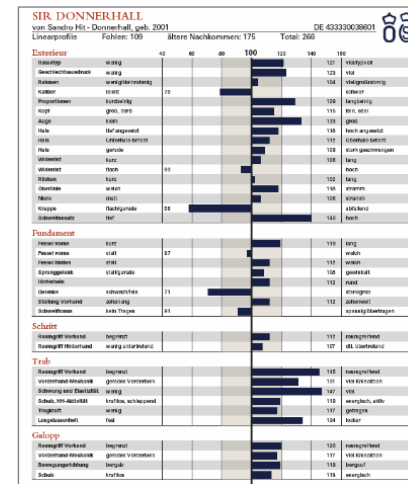
Routine genetic evaluation for linear traits in the Oldenburg studbooks (OL, OS)

- ✓ since 2017, annually in October / November
- ✓ conformation and performance (46 traits)



Genetic linear profiles: content

- relative breeding values
 - mean of 100, genetic standard deviation of 20 (orientation as in the linear scheme)
 - base definition: sires born after 1994 with at least 3 linearly described adult progeny (GE 2020: N=527 sires)
- genetic stallion profiles: two groups



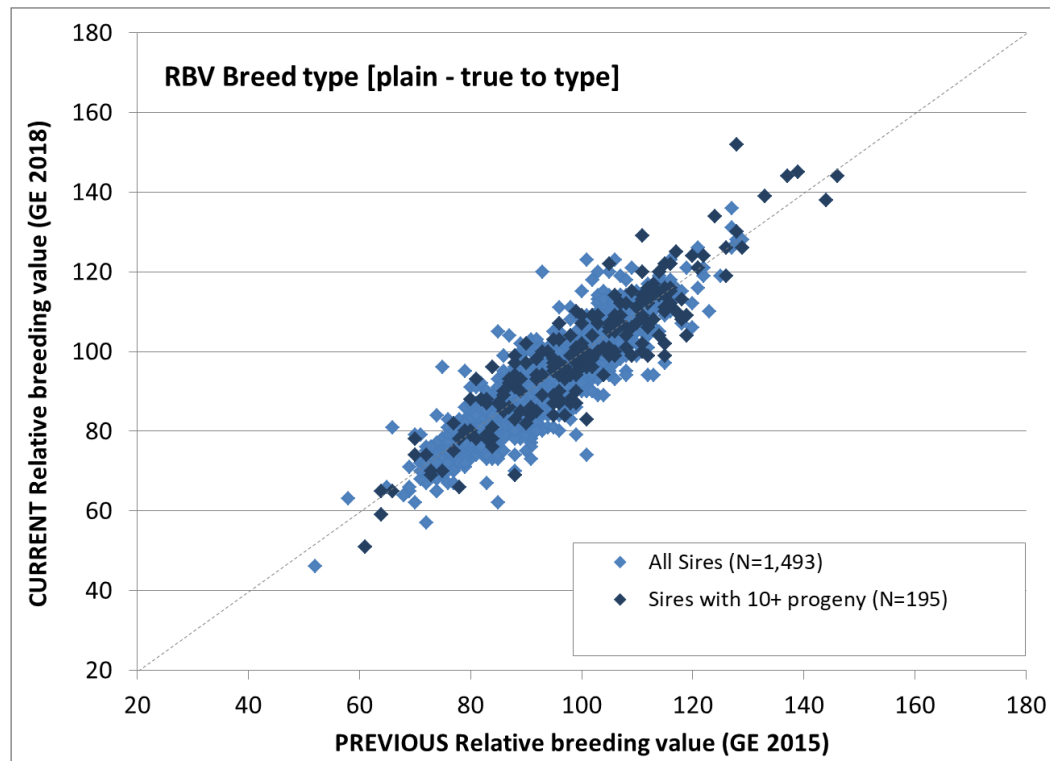
- (1) frequently used sires (≥ 20 linearly described progeny; age ≥ 7 years)
 → index as weighted combination of EBV: $40\% EBV_{Foal} + 60\% EBV_{Adult}$
- (2) young sires (maximum age of 6 years, minimum of 8 foals)
 → EBV for linear traits in foals, i.e. EBV_{Foal}

Genetic linear profiles: use

- expectations?
 - reliable 'filtering out' of genetic dispositions (better than phenotype-based progeny statistics)
 - earlier, more objective and more helpful information for breeders
- concrete questions?
 - Can we use genetic profiles of young stallions with few linearly described foals (first crop) for support of mating decisions?
 - Will the genetic linear profiles change over time? If so, how much?
 - Can we predict the progeny phenotypes by the genetic linear profiles of their fathers? How well?



Changes over time? Example I

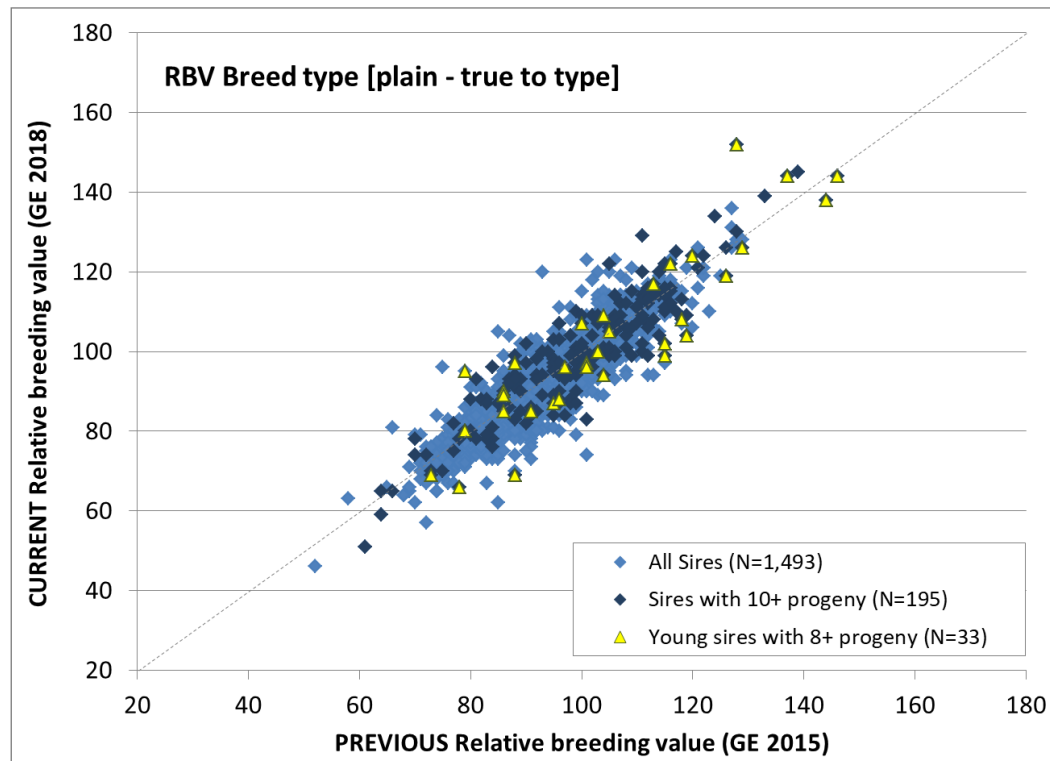


Explanation:

comparison of results from GE test run with truncated data (GE 2015) with regular GE run using all data (GE 2018) for:

- all sires with progeny already in GE 2015 (on average only 6 progeny)
- sires with more reliable EBV (10 or more progeny already in GE 2015)

Changes over time? Example I

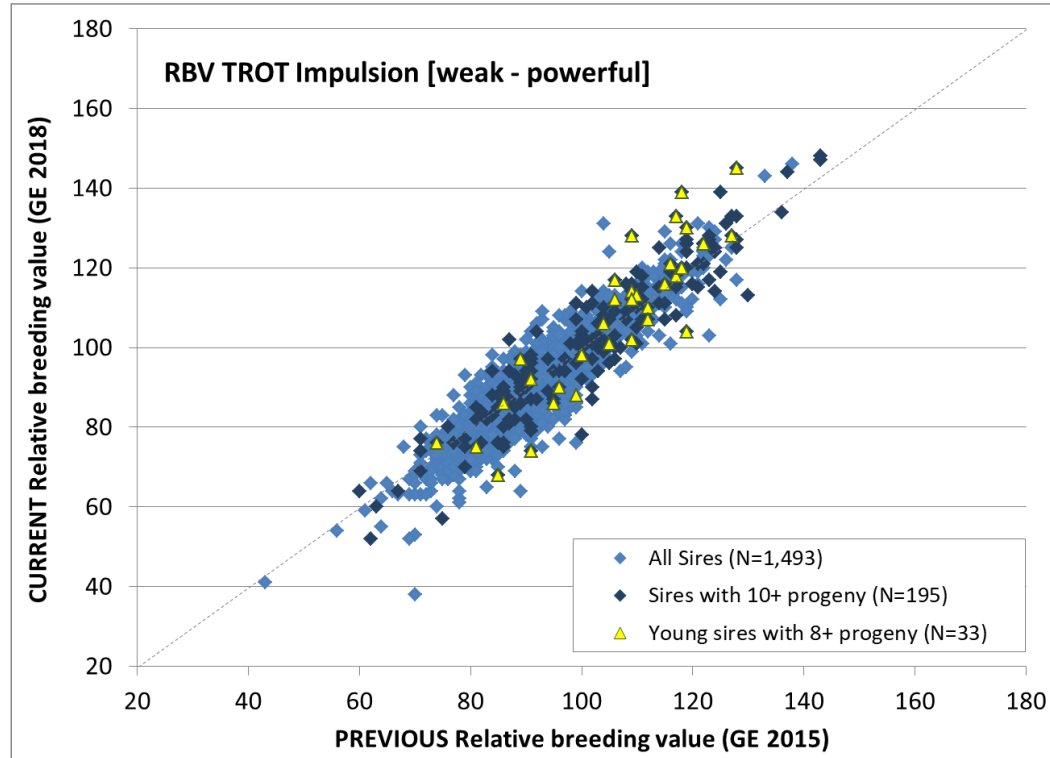


Explanation:

comparison of results from GE test run with truncated data (GE 2015) with regular GE run using all data (GE 2018) for:

- all sires with progeny already in GE 2015 (on average only 6 progeny)
- sires with more reliable EBV (10 or more progeny already in GE 2015)
- young sires (max. age of 6 years) in GE 2015

Changes over time? Example II



Explanation:

comparison of results from GE test run with truncated data (GE 2015) with regular GE run using all data (GE 2018) for:

- all sires with progeny already in GE 2015 (on average only 6 progeny)
- sires with more reliable EBV (10 or more progeny already in GE 2015)
- young sires (max. age of 6 years) in GE 2015



Study answers to breeders' questions

- *Can we use genetic profiles of young stallions with few linearly described foals (first crop) for support of mating decisions? **YES – they are valuable early indicators.***
- *Will the genetic linear profiles change over time? If so, how much? **YES – changes are possible and expected, can be substantial.***
more progeny with linear data → reliability of breeding values ↑ → changes ↓
- *Can we predict the progeny phenotypes by the genetic linear profiles of their fathers? **How well?***
YES – prediction is possible, advanced use of linear data implies continuous improvement.

- support of expectations regarding linear profiling
 - reliable 'filtering out' of genetic dispositions
(better than phenotype-based progeny statistics)
 - earlier, more objective and more helpful information for breeders



Conclusions & perspectives |

- successful routine implementation of linear profiling as important step towards more sustainable competitive breeding
 - substantial gain in information on conformation and performance
 - increased transparency
 - feasible tool for collection of high quality phenotypic data required to benefit from more targeted breeding applications
- **to be accompanied by continuous data quality management for optimum use of linear description**



Conclusions & perspectives II

- successful routine implementation of linear profiling as important step towards more sustainable competitive breeding
- high value of refined linear phenotypes
 - specific individual characterization (linear profiles of presented horses)
 - improved genetic characterization (genetic stallion profiles)
 - appropriate basis of future genomic applications (genomic profiles)
- **engagement for linear profiling = investment in the future**

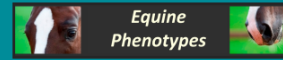




IT-Solutions for
Animal Production

Contact: PD Dr. habil. Kathrin F. Stock
(E-mail: friederike.katharina.stock@vit.de; phone: +49-4231-955623)

Further information on
linear profiling and related activities:



TAKE HOME:

- **engagement for linear profiling = investment in the future**
- new and improved phenotypes as suitable targets for new and improved breeding applications
- collaborative approaches to capture the full potential using genetic and genomic tools

Thank you !