62nd annual meeting of the EAAP 2011, Stavanger / Norway

S21_8 (abstract no. 10894)

Genetic analyses of movement traits in German Warmblood horses

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Background

- movement evaluations as integral part of selection processes in the Warmblood horse
- remarkable breeding progress in key features of riding horse performance with regard to the gaits (e.g., impulsion, ground cover and expression in trot)
- unfavorable movement characteristics as long-known phenomenon with rare, but regular occurrence in horses of different age
 - \rightarrow sparse knowledge about quantitative and qualitative importance, causative factors, ...

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Information on movement

- routine gait evaluations
 - regular breeding events
 - free movement, movement under rider
 - subjective scores (1-10)
- detailed movement evaluations
 - regular breeding events
 - (primarily) free movement
 - documentation of specific favorable and unfavorable movement characteristics

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Study approach



- detailed movement evaluations from regular breeding events of the Oldenburg horse breeding societies (OLD, OS) in 2009 and 2010
 - foal registrations (foals and mares)
 - mare shows (mares)
- routine gait evaluations
 - studbook inspections (SBI) of OLD+OS in 2009
 - mare performance tests (MPT) of OLD+OS in 2000-2008



genetic correlation analyses

to learn more about the role of unfavorable movement characteristics seen in juvenile and adult horses

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Detailed movement evaluations

- foals (n=3,374)
 - mean age of 2.3 months
 - free movement at the side of their dams
 - descending from 476 sires with 1-137 (Ø 7.1) offspring
- mares (n=2,844)
 - mean of 9.2 years (3.2 years in the show mares, 10.8 in the active broodmares)
 - presented in walk and trot at hand
 - descending from 1,197 sires with 1-126 (Ø 3.5) offspring

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Movement traits

- new movement traits from detailed movement evaluations
 - irregular tail tone and/or posture (TTP)
 - irregular motion pattern in hind legs
 - irregularity in general motion pattern
 - pace in foals

indications of imbalance (IMB)

- standard traits from SBI and MPT
 - walk, impetus & elasticity, correctness of gaits
 - walk, trot and canter during free movement, walk, trot and canter under rider, rideability, free jumping

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Distributions and heritabilities

Movement trait	Foals (n=3,374)		Mares (n=2,844)		
	Prev.	$h^2 \pm SE_{h^2}$	Prev.	$h^2 \pm SE_{h^2}$	
TTP (irregular tail tone and/or posture)	3.97%	0.076 ± 0.030 (0.395 ± 0.158)	4.89%	$0.043 \pm 0.019 \ (0.194 \pm 0.088)$	
IMB (indications of imbalance)	6.22%	0.117 ± 0.035 (0.457 \pm 0.137)	5.49%	0.052 ± 0.022 (0.220 \pm 0.094)	

results from bivariate analyses of corresponding traits in foals and mares with heritabilities on observed scale (first line) and underlying liability scale (second line)

SBI trait (n=1,987)	Mean ± SD (range)		
Walk	6.60 ± 0.67 (5.0-9.0)		
Impetus & elasticity	6.58 ± 0.74 (5.0-9.0)		
Correctness of gaits	6.22 ± 0.67 (3.0-8.0)		

 h^2 =0.3-0.6 for most SBI and MPT traits (exceptions: Correctness of gaits h^2 =0.1, and rideability h^2 =0.2)

MPT trait	Mean ± SD (range)			
(n=2,758)	Free	Under rider		
Walk	7.23 ± 0.57 (5.5-9.0)	7.24 ± 0.64 (5.0-10.0)		
Trot	7.39 ± 0.60 (5.5-9.5)	7.11 ± 0.69 (5.0-10.0)		
Canter 7.24 ± 0.54 (5.5-9.0)		7.34 ± 0.61 (5.0-9.5)		
Rideability		7.38 ± 0.67 (5.0-9.5)		
Free jumping	7.13 ± 0.82 (4.5-10.0)			
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Genetic correlation analyses (I)

Estimation of genetic parameters

- Residual Maximum Likelihood (REML) / VCE6
- · bivariate linear animal models
 - movement trait (binary)
 - SBI or MPT trait (quasi-continuous)
- → additive genetic correlations (r_g)

Prediction of breeding values

- Best Linear Unbiased Prediction (BLUP) / PEST
- univariate linear animal models
- → Pearson correlation coefficients (r_{EBV})

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Genetic correlation analyses (II)

Pedigree data

- · unified animal ownership database (vit)
- relationship matrix with 44,158 horses

Models

MoveF, MoveM

$$y_{iklpst} = \mu + b*age_m_i + BMON_k + JUDGE_l + eventM_p + a_s + e_t$$

 $y_{jklmpst} = \mu + b*age_j_i + BMON_k + JUDGE_l + ETYPE_m + eventM_p + a_s + e_t$

SBI, MPT

$$y_{knqst} = \mu + BMON_k + SBI_AGE_n + eventSBI_q + a_s + e_t$$

 $y_{korst} = \mu + BMON_k + MPT_AGE_o + eventMPT_r + a_s + e_t$

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Genetic correlations

Additive genetic correlations with $\[\] r_g < -0.1 \]$ (favorable) $\[\] r_g > 0.1 \]$ (unfavorable) and Pearson correlations between breeding values (r_{EBV}) of sires*

SBI or MPT trait	Foals		Mares	
	TTP	IMB	TTP	IMB
Walk	0.092	0.048	0.338	0.312
Impetus & elasticity	0.054	0.032	0.495	0.450
Correctness of gaits	0.004	0.103	0.239	0.211
Walk - free	0.105	0.133	0.314	0.275
- under rider	0.155	0.088	0.449	0.427
Trot - free	-0.042	0.006	0.310	0.320
- under rider	-0.018	0.095	0.220	0.196
Canter - free	-0.166	-0.075	0.165	0.181
- under rider	-0.103	-0.079	0.262	0.275
Rideability	-0.044	0.004	0.344	0.366
Free jumping	-0.106	-0.176	-0.315	-0.308

TTP = irregular tail tone and/or posture; IMB = indications of imbalance

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^{*} r_{EBV} for sires with ≥5 offspring with detailed movement evaluations or SBI / MPT and offspring in each of the trait groups (MoveF, MoveM, SBI: n=94; MoveF, MoveM, MPT: n=73)

Conclusions

- genetic variability in the new movement traits (TTP, IMB)
 reflecting unfavorable movement characteristics:
 possible use of detailed movement information (foals > mares)
 to select for better balance in movement
- indications of unfavorable genetic correlations between the new movement traits and standard traits from SBI and MPT:
 - SBI/MPT traits
 no measures of breeding progress with regard to specific movement characteristics (balance)
 - TTP, IMB
 no reduction of unfavorable movement characteristics through
 selection based on subjective scores from routine gait evaluations

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Implications

- need for deeper understanding of unfavorable movement characteristics, particularly in adult horses (refined recording including evaluation conditions)
- · benefits from systematic use of foal data
- unfavorable movement characteristics as another argument for revising the traditional scoring system in conformation and performance evaluations for breeding purposes
 - → linear scoring reflecting (favorable and unfavorable) specific movement characteristics

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