

Federation of Animal Science, Ghent, Belgium





# Importance of high quality data base systems in the study of genetic characteristics in horses

M. Wobbe<sup>1,2</sup>, K. F. Stock<sup>1,2</sup>, F. Reinhardt<sup>1</sup>, J. Tetens<sup>3</sup>, R. Reents<sup>1</sup>

<sup>1</sup>IT Solutions for Animal Production (vit), Verden, Germany; <sup>2</sup>University of Veterinary Medicine Hannover (Foundation), Hanover, Germany; <sup>3</sup>University of Goettingen, Goettingen, Germany

#### **Outline**



- role of genetic defects and characteristics
  - genetic background and examples (cattle, horse)
- role of data base systems: monitoring and interpretation
- Warmblood Fragile Foal Syndrome (WFFS) as a famous example
  - disease characteristics
  - statistical analyses: WFFS mutation and equine reproduction data
  - distribution patterns
- conclusions



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#### Genetic background

- replication error during meiosis (parent  $\rightarrow$  offspring)
  - apparent in the next generation
  - consequence could be: neutral, trait, disease or unknown
  - often recessively inherited
- ❖ selection → increase of inbreeding
  - stronger relevance of mutations
- extensive recording of phenotypes, ancestral and genomic information
  - facilitating the discovery of mutations



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### **Example: Cattle breeding**

- since introduction of genomics
  - discovery of several deleterious genetic defects (BLAD, Brachyspina etc.)
  - as well as: single gene effects without any negative consequences (colour genotypes, polledness, casein etc.)
- → genetic <u>characteristics</u> instead of genetic defects



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#### **Genetic characteristics in horses**

- routine screening in some horse breeds, e.g.:
  - Arabians (SCID, CA, LFS etc.)
  - Quarter Horses (PSSM, HERDA, HYPP etc.)
- in the past: Warmblood breeders not so familiar with molecular genetic testing
- main point of contact with laboratory work:
  - parentage testing
  - testing for colour genotypes





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### Role of data base systems



- how to control, understand and discover genetic characteristics?
  - > informative and enough data
  - > stored and managed in one central location
- comprehensive data base system needed for:
  - monitoring and interpretation of genetic characteristics
    - essential for a responsible handling by the breeding organizations



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- hereditary connective tissue disorder in Thoroughbred and Warmblood breeds
- first described in 2011 (US research group)
  - commercial genetic test since 2013
- became widely known among breeders in 2018
  - ➤ WFFS case in the USA → discussed internationally
- → WFFS as an example how to elucidate a genetic characteristic with the help of a comprehensive data base system



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#### WFFS – disease characteristics



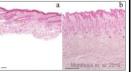
- point mutation in the PLOD1 gene, recessively inherited
- consequences of mutation:
  - thin and fragile epidermis not firmly attached to the subcutaneous tissue
  - even low stress leads to skin rupture
  - hyperextensible joints of the limbs
- ❖ only few affected (homozygous) foals are born → not viable
- → what about WFFS related losses?











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## Hardy-Weinberg equilibrium (HWE)



 null hypothesis of relationship between allele and genotype frequencies



- reported carrier (heterozygotes) frequency of 9.5-15 %
- mating a carrier sire to the average mare population
  - estimated value of 2.4-3.7 % homozygous offspring (following HWE)

→ 2.4-3.7 % more foal losses (because of WFFS) expected

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# **HWE – Explanation**



probabilities of genotypes with random mating (11 % carrier freq.)

| mare pop.    | AA (89%)                   | Aa (11%)                                   |
|--------------|----------------------------|--|
| AA (free)    | all free (AA)              | ½ free (AA)<br>½ carrier (Aa)              |
| Aa (carrier) | ½ free (AA) ½ carrier (Aa) | ¼ free (AA) ½ carrier (Aa) ¼ affected (aa) |

 $0.25*0.11 = 0.028 \leftarrow$ 

expected: 2.8 % homozygous offspring (more foal losses)

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## Analyses with equine reproduction data



- covering data from 10 German Horse Breeding Associations
  - N = 426,568 coverings of 10 years (2008-2017)
  - N = 177,582 coverings from stallions with known WFFS status
- analyses of variance with SAS software (v. 9.2, PROC HPMIXED)
  - dependent variable (1/0): foal was born (1) or foal was not born / died within the first 2 days (0)
  - <u>fixed effects</u>: WFFS status of the sire, data provider, covering year, age of the mare
  - <u>random effect</u> of the sire himself + random residual

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## Analysis of variance - Results



- expectation (following HWE):
  - ≥ 2.4-3.7 % more foal losses among carriers than free sires
- results based on all sires with known WFFS status
  - least square means of the foaling rates

| dataset restriction          | least square means foaling rates |        | difference  |                   |
|------------------------------|----------------------------------|--------|-------------|-------------------|
|                              | carrier                          | free   | free vs. ca | rrier WFFS status |
| none (a)                     | 0.6393                           | 0.6636 | 0.0243      | 0.0304            |
| min. 5 foals per sire (b)    | 0.6388                           | 0.6638 | 0.0250      | 0.0264            |
| min. 5 sire per provider (c) | 0.6550                           | 0.6783 | 0.0233      | 0.0397            |
| combination of (b) and (c)   | 0.6552                           | 0.6792 | 0.0240      | 0.0347            |

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## WFFS – Distribution patterns (I)

- WFFS known for several breeding populations (cases in USA, SWE, CH, DK, GER)
  - mutation must be rather old
  - > first occurrence in a founder who is an ancestor in many populations
  - → English Thoroughbred was / is used in many different breeds
- data analyses: WFFS tested horses (N = 3576) and their ancestry
  - → which founder appears in the pedigree of all known carriers?

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# WFFS – Distribution patterns (II)



- only horses which appear in every known carrier
  - Dark Ronald xx, born 1905
  - his father Bay Ronald xx, born 1893
  - → very influential English Thoroughbred horses
- skeleton, heart + skin of Dark Ronald xx in a museum in Halle (GER)
  - > sample from heart / skin was taken
  - DNA extraction and testing in progress
  - ightarrow spread of mutation i.a. via Dark Ronald xx and others
- note: origin of the mutation most likely much older



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#### **Conclusions**



- inheritance theory confirmed by foaling rates
  - WFFS as a cause of premature foal losses
- to avoid affected / stillborn foals and abortion: use the possibilities of modern breeding!
  - WFFS can be controlled by targeted testing and mating
- one distribution path of the mutation could be found
  - → not possible without the power of strong data base systems

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