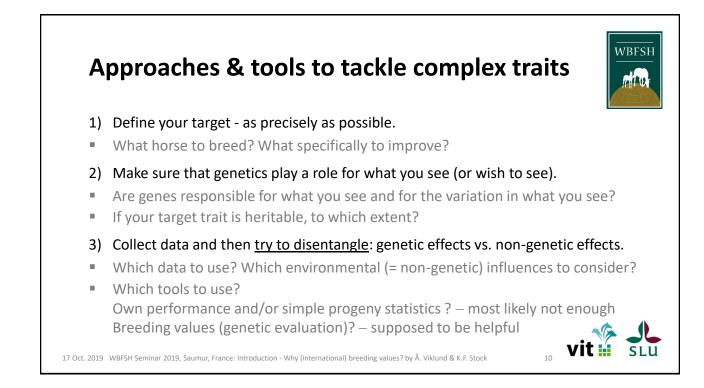
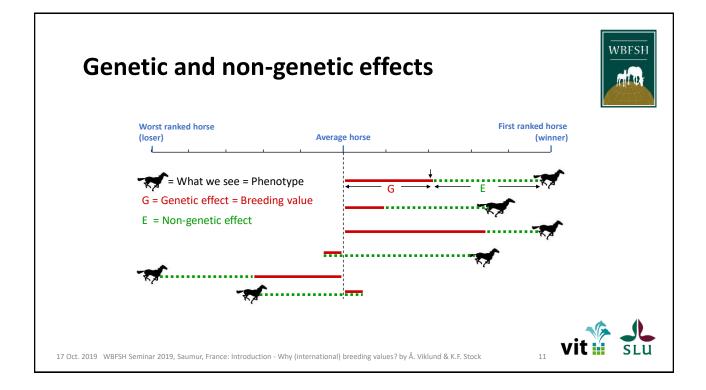


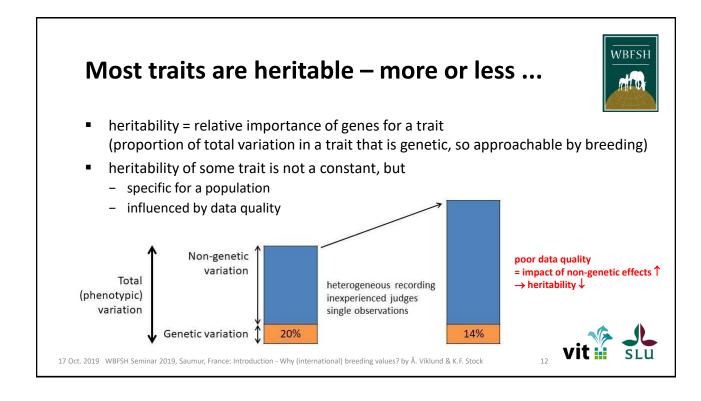


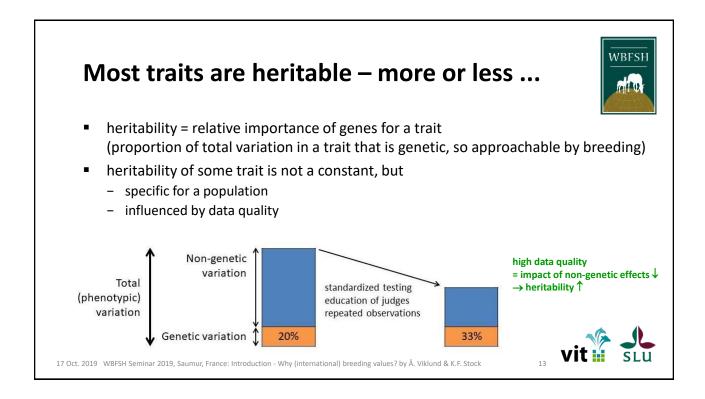
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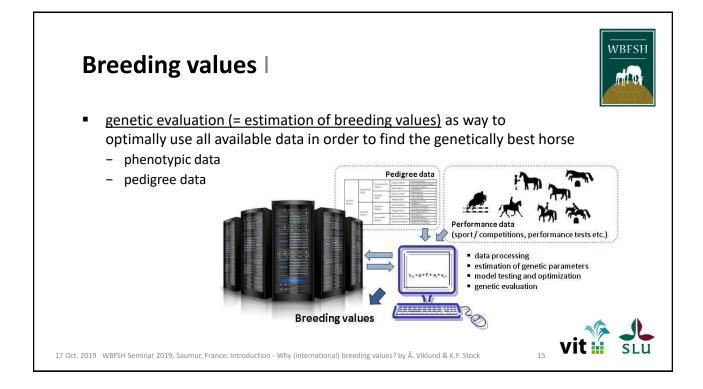
# Heritability & breeding tools

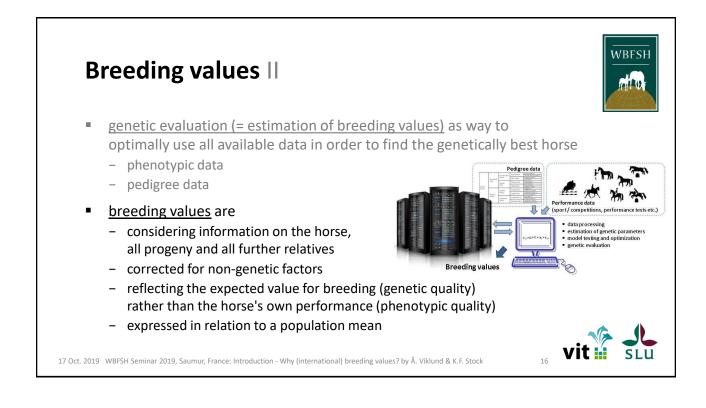


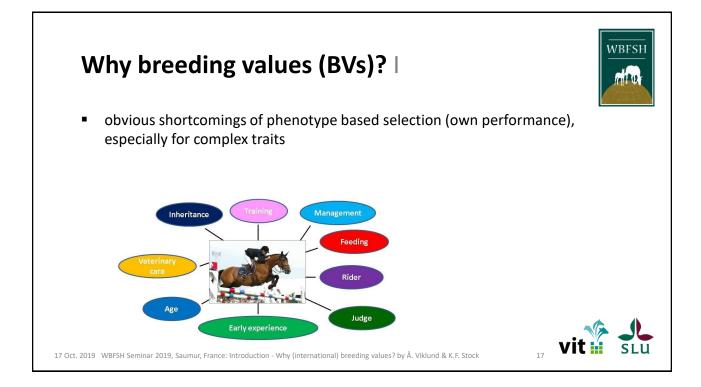
14

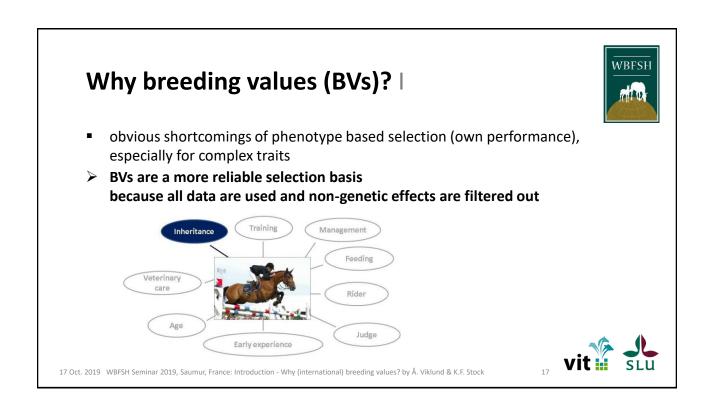
- highly heritable traits
  - $\rightarrow$  good chances to achieve reasonable breeding progress through 'breeding by eye'
- traits of low heritability
  → need for tools to find the best horses not only by chance

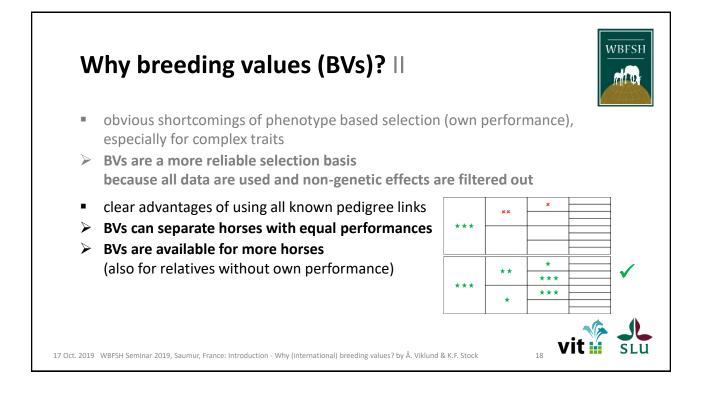
Heritability level	Examples of traits	Amount of data needed	Approach and methodology / tools
high	basic morphology aspects like size (withers height) with $h^2\approx 0.45\text{-}0.55$	+	easy e.g. phenotypes
moderate	conformation traits, some performance traits like certain gait characteristics with $h^2 \approx 0.20$ -0.35	++	advanced e.g. breeding values
low	many performance traits, behavior, health traits with $h^2\approx 0.10$ or less	+++	sophisticated e.g. (genomic) breeding values

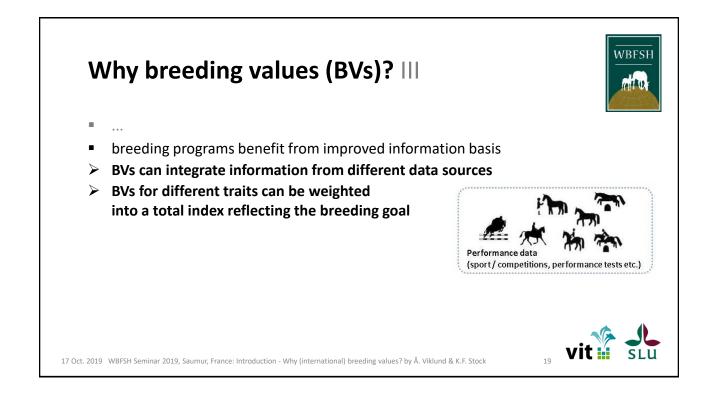


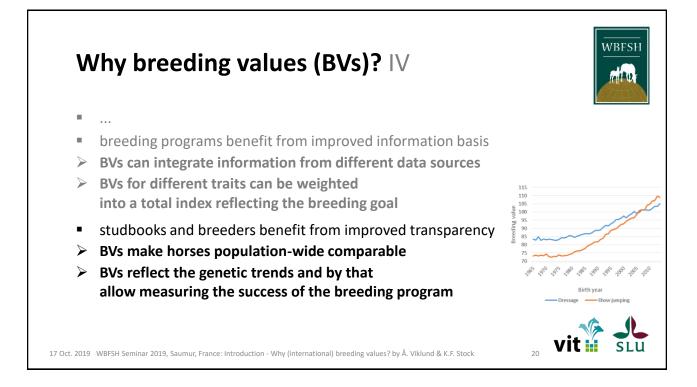


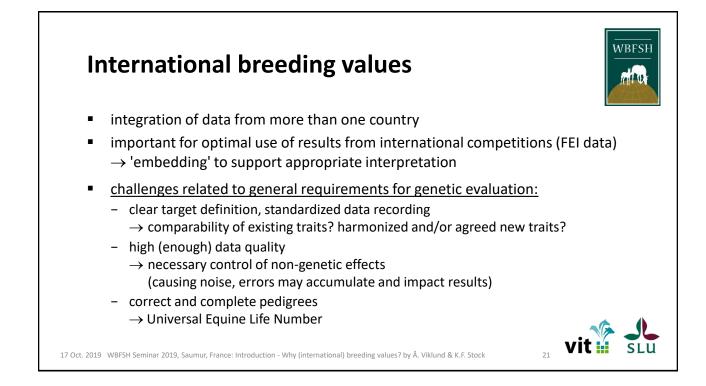










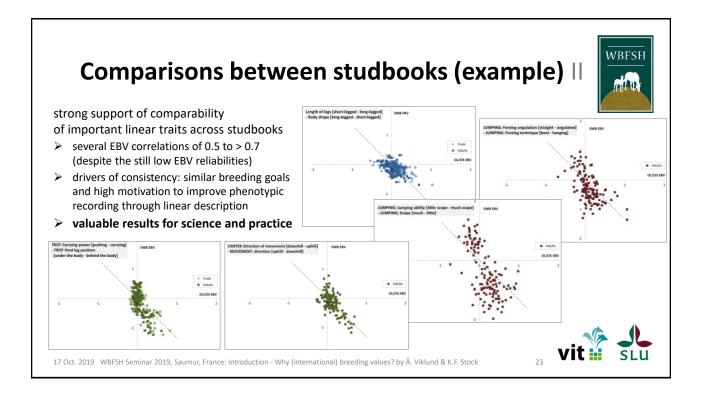


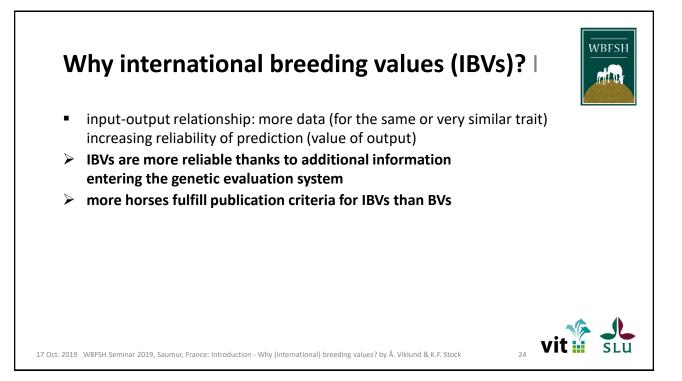


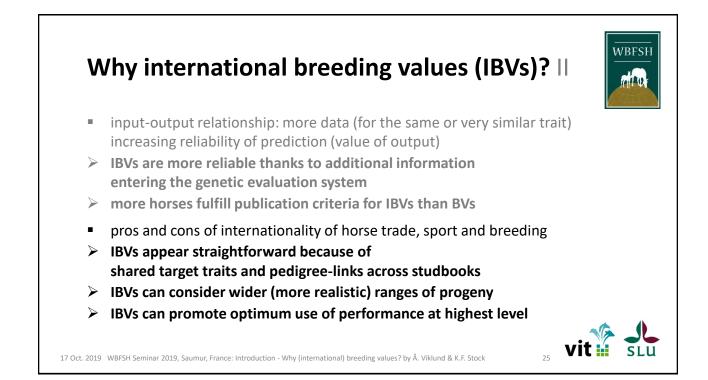
- pilot study in connection with new applications for linear profiling
- ongoing collaboration between studbooks:
  Oldenburg (OL, OS) and Swedish Warmblood (SWB)
- use of breeding values of stallions for linear traits

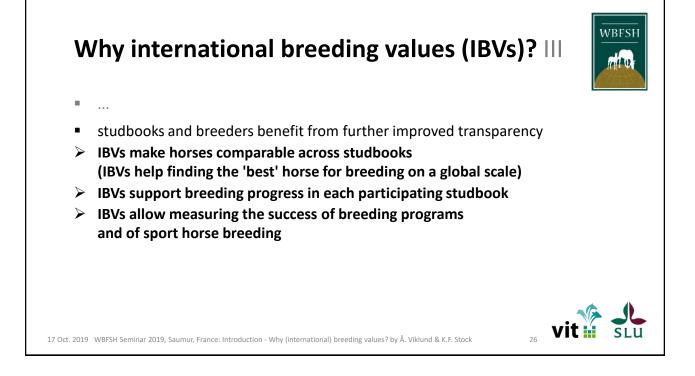
Data structure with reference to figures from the first presentation of study results (prototypes of genetic evaluations in 2016).

Sire group characteristics	Progeny group	No. of linearly described progeny per sire within studbook		
Sile group characteristics		OL/OS (N <sub>sires</sub> =1,627)	SWB (N <sub>sires</sub> =331)	
All stallions:	foals	7.7 (max. 181) / N <sub>sires</sub> =864	-	
linearly described progeny in OL/OS or SWB	≥2.5-year-olds	3.4 (max. 101) / N <sub>sires</sub> =1,114	5.7 (max. 69)	
Stallion sample (N=132):	foals	18.2 (max. 174) / N <sub>sires</sub> =87	-	
linearly described progeny in OL/OS and SWB	≥2.5-year-olds	11.9 (max. 101) / N <sub>sires</sub> =119	4.4 (max. 46)	









## **Prospects**



- great potential of international breeding values for WBFSH member studbooks and for sport horse breeding as a whole
- clearly defined tasks
  - → different starting points, but options to work towards IBVs for all studbooks

# Thank you!

### What is needed?

- well defined target what do we want to achieve?
- systematic and high quality data recording
  - traits reflecting the breeding goal
  - standardized recording
  - many horses, representative for the population
- correct and complete pedigree data
- reliable infrastructure: geneticists and computers <sup>(2)</sup>
- challenges of the IBV project to be met by collaboration making best use of available resources and expertise

