



How international collaboration fostered an efficient use of the genomics for a reliable cattle breeding INTERBULL: INDUSTRY SEMINAR IN VERDEN (GERMANY) 25/02/15

Clotilde Patry, EuroGenomics

Contents

EuroGenomics key figures

- How breeding companies and farmers took up genomic tools
- Joined efforts to keep improving the quality of the tools



Presentation of EuroGenomics





Since 2009: 5 then 7 members joined forces for a reliable cattle

9 European countries



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Genomika Polska

Interbull Industry Seminar, Verden, Germany - 25/02/15

VIKINGGENETICS

Presentation of EuroGenomics





Since 2009: 5 then 7 members joined forces for a reliable cattle 9 European countries 15 millions of AI in Holstein breed

Merge of the reference population

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About 30,000 candidates (before GS: < 3000 progeny tested bulls)

Official information for about 1,500 young bulls per year (from 10 months)



 $\frac{3}{4}$ females $\frac{1}{4}$ males 9% 54k Females 69% EuroG10k Females

EuroG10k Males

~130,000 Holstein animals:

54k Males

Contrasted strategies between countries

Distribution in 2014 (beeing adapted over the years)



Accelerated genetic progress



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A quick adoption of the genomics in the breeding schemes



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How to keep improving the quality of the GEBV?





A better use of the available information

- From the reference population
 - Harmonisation of the traits
 Quality of the prediction equations



More relevant information for decisions on selection



- EuroG10k = a customized chip, adapted to the needs, quickly evolving (v4)
- A result from collaboration (research & industry)
- Increased the quantity of relevant information :
 - Imputation
 - Genetic abnormalities
 - Genes of interests

- Quality of the predictions
- Mating plan, breeding schemes



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More information at the time of Al use

« Domestic » versus « foreign » bulls : reliability of the predictions should be the same with genomics

- > 2014: Rankings of young bulls also became international
 - Exchange of genotypes
 - Participation to GMACE
- Which encouraged
 - A large number of very young bulls to be published
 - \Rightarrow with or without semen available
 - A quick turnover while breeding values are expected to evolve (R²)

More information at the time of Al use

« Domestic » versus « foreign » bulls : reliability of the predictions should be the same with genomics

- ► The most complete information for the computation of GEBV
 - Exchange of genotypes of the young bulls
 - Completeness of the relationship information (genomic and non genomic data)
 - \Rightarrow About 5000 additional bulls (/ Reference Population)
- Result: early information for each YB on different scales
- \Rightarrow Leverage to make the methodologies evolve
- \Rightarrow Useful for users: within the EG area and outside the EG area

More information at the time of Al use

For farmers outside of the EG area (without own genomic evaluations): increased reliability



Conclusions

- A quick adoption of the GEBV by breeding companies and by farmers
- Various needs of the farmers, various strategies and practices between countries
- Reliability level are still expected to be increased
- Joint efforts: research, calculation centers and industry accepted to collaborate in the interests of the dairy farmers
- A new way of collaboration = co-opetition



