

International issues in dairy cattle genetics

Vincent Ducrocq

Jouy-en-Josas, France



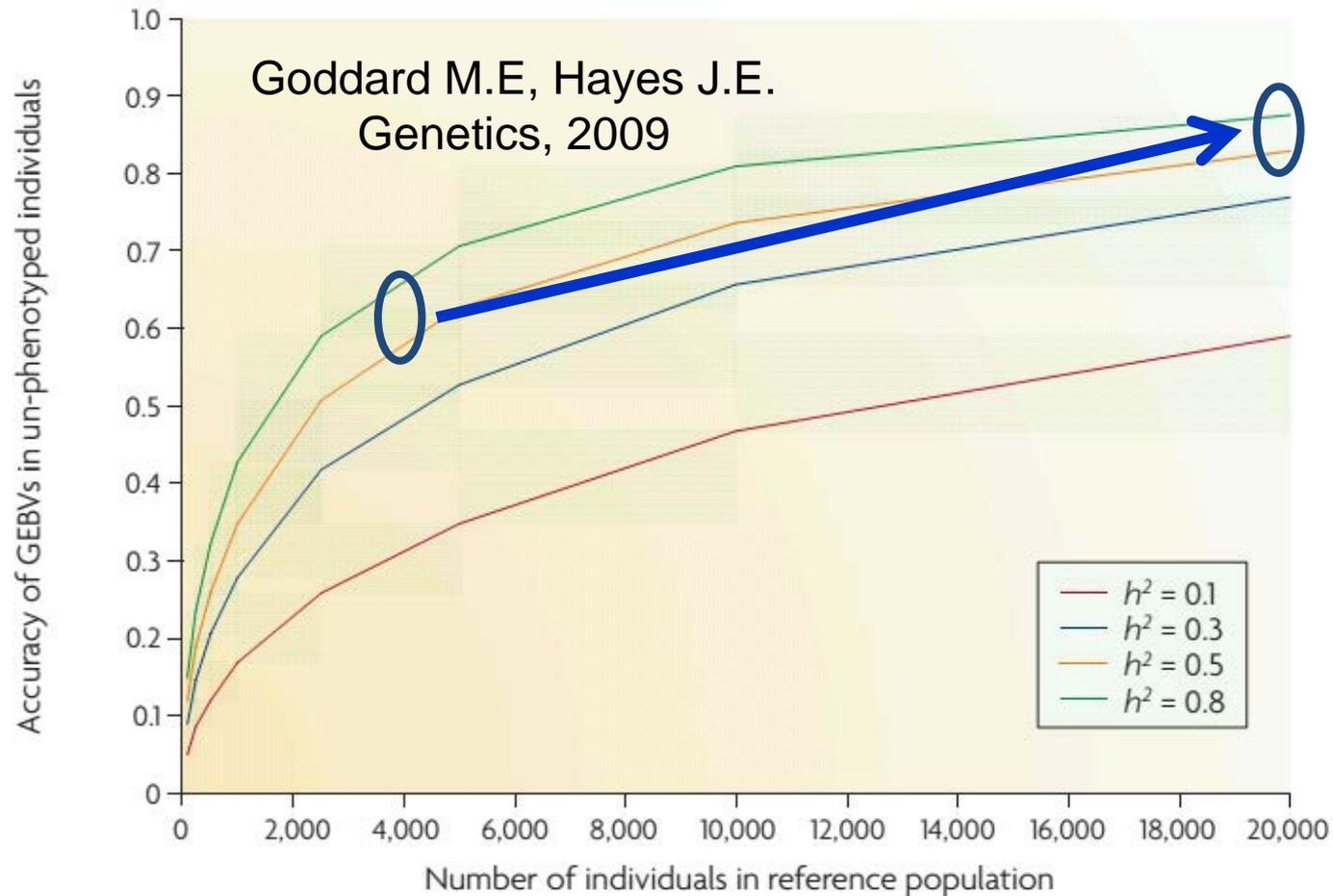
Outline

- New context: **genomic era**
- Principles of **international genetic evaluations / Interbull**
- How do farmers get access to the best bulls with genetic or genomic evaluations
 - *from countries which are Interbull members*
 - *from the others (with/without genetic or genomic evaluations)*
- Key messages

Genomic selection: very fast adoption

- Started between 2008 and 2010
- **Reference population** = genotyped progeny-tested bulls from which estimates of the effect of tiny chromosome segments are computed
- Reference population size = key parameter of accuracy of genomic selection

Importance of reference population size



Genomic selection: very fast adoption

- Tens of thousands of young calves genotyped each year
- Organized progeny testing: decreased or abandoned
- **Market share of semen from young bulls: 40 to 75%**

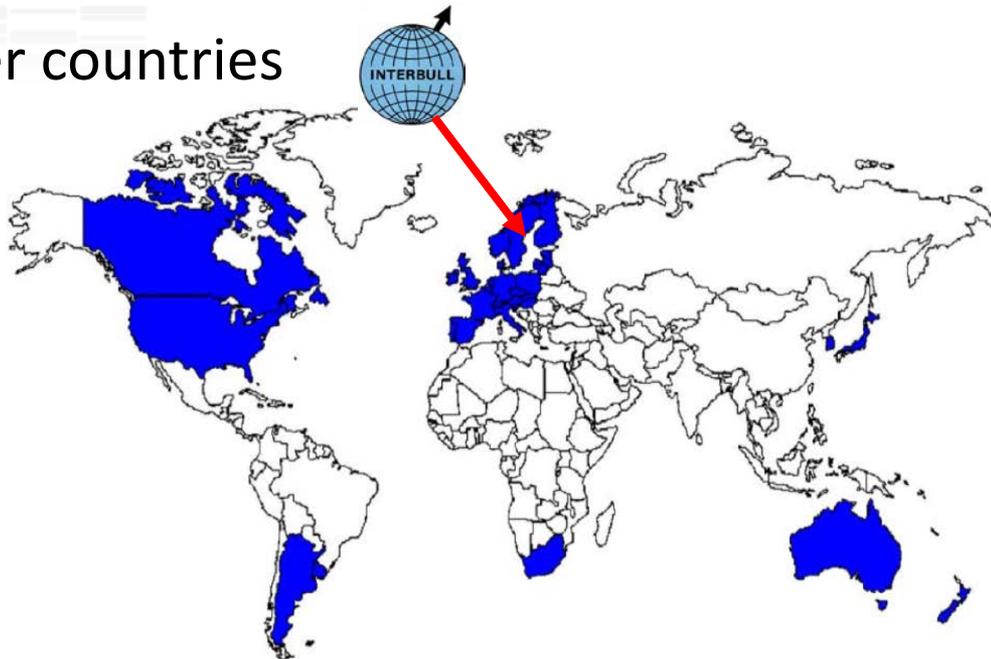
International Genetic Evaluations

- National evaluations are not directly comparable (different traits, different production system and environments, different recording systems, different models of analysis)
- For the past 30 years, **Interbull** has made such comparisons possible (+ has contributed to the international recognition of national evaluations (harmonization, validation))



Interbull

- 32 member countries

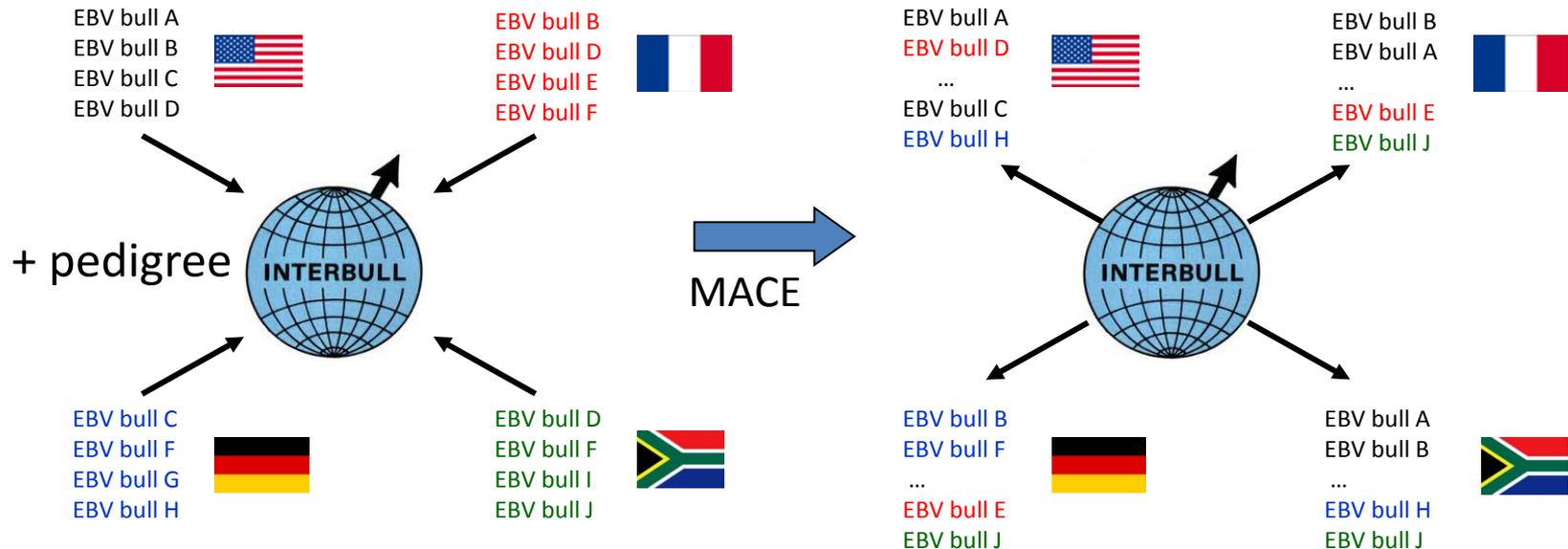


- 6 (groups of) breeds: Holstein, Simmental (including Montbéliarde) Brown Swiss, “Red” Dairy cattle, Jersey, Guernsey.
- Traits: Production, Type, Udder Health, Fertility, Longevity, Workability, Calving traits

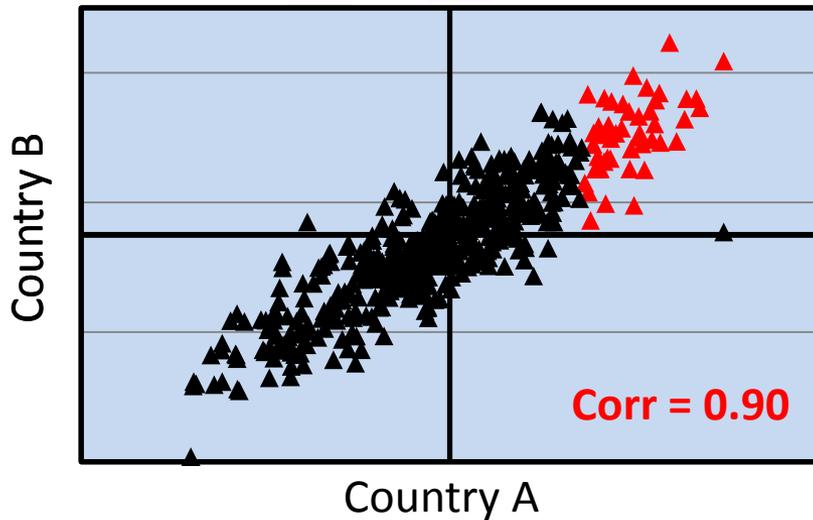
International genetic evaluations

➤ Meta-analyses of national results of bulls

MACE = Multiple Across Country Evaluation



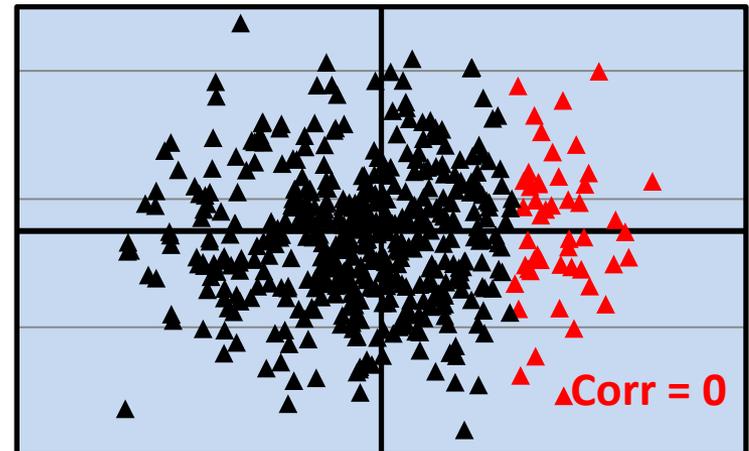
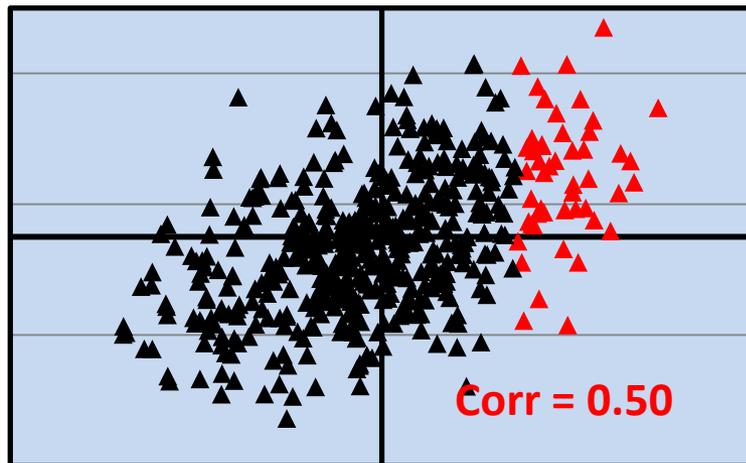
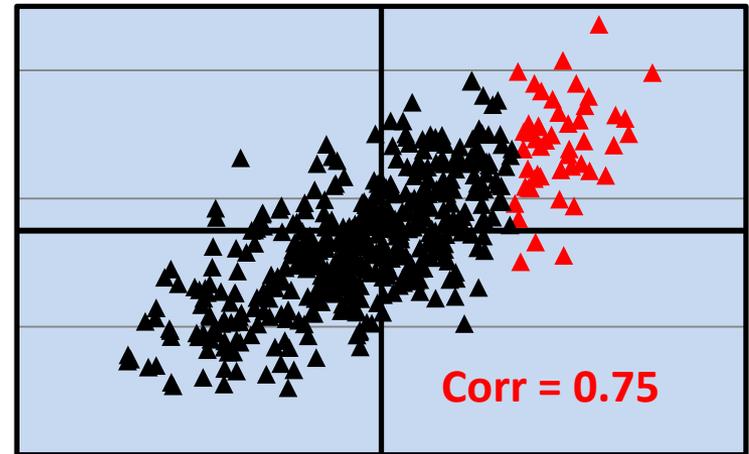
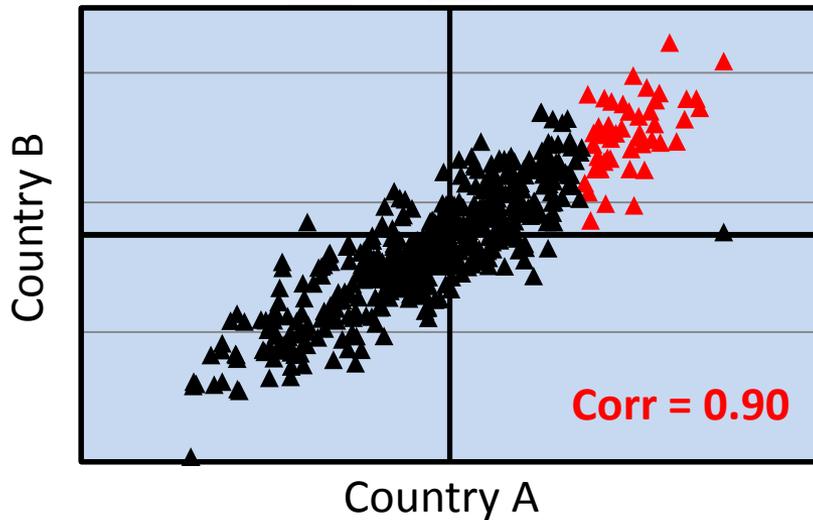
How much reranking ?



Within Europe or North America
(for production traits)

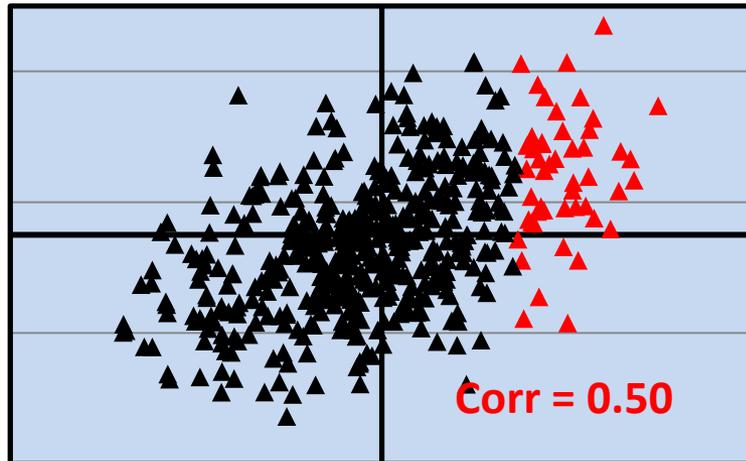
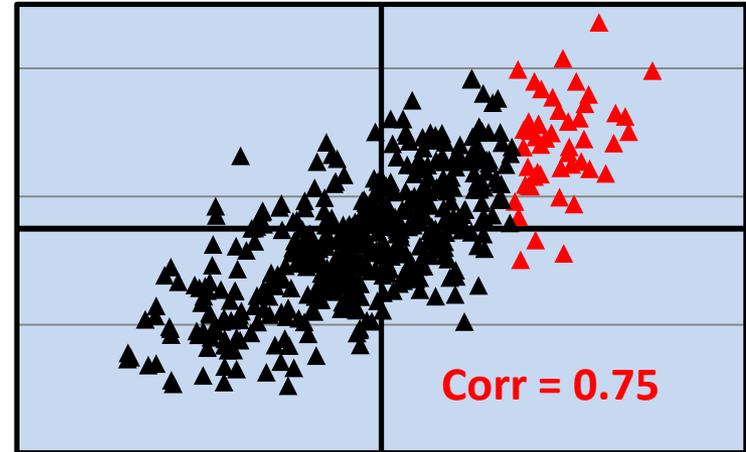
Depends on **genetic correlation**
between countries
(= a measure of Genotype x
Environment interaction)

How much reranking ?



How much reranking ?

Between New Zealand or
Ireland and continental Europe
(or some fertility traits)



Between Kenya and UK
(Ojango et al, 2003)

Farmers /breeding organisations: Consequences

➤ Strongly depends on the country situation

- ✓ Is it an Interbull member? (access to international ranking on own national scale)

➤ Even more complex with genomic evaluations

- ✓ Access to a (inter)national reference population ?

Interbull members

Top 10 LIST:

1. _____
2. _____
3. _____
4. _____
5. _____
6. _____



Top 10 LIST:

1. _____
2. _____
3. _____
4. _____
5. _____
6. _____



Top 10 LIST:

1. _____
2. _____
3. _____
4. _____
5. _____
6. _____

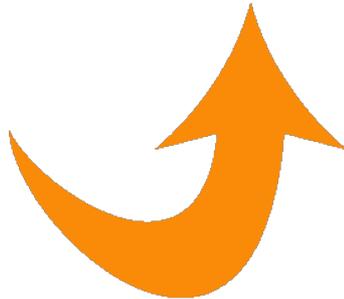


TOP 100



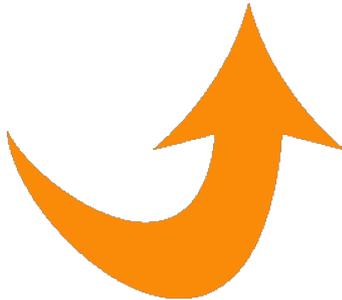
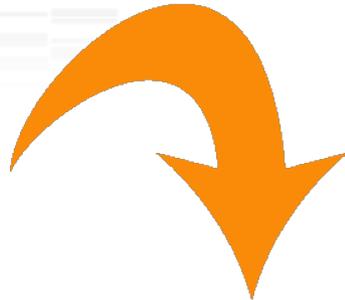
Top 10 LIST:

1. _____
2. _____
3. _____
4. _____
5. _____
6. _____



Non members

✓ Europe? too many scales !!!
which one to choose ?



US scale

	Net merit (NM\$) (2010)	
(Milk)+Fat+Protein	35	
Longevity (PL)	22	} 48
Somatic cells	-10	
Fertility (DPR)	11	
Calving diff./ Stillbirth	-5	
Type composite		
Udder composite	7	} 17
Feet/legs composite	4	
Body size	-6	



Source 1: Net merit as a measure of lifetime profit: 2010 revision *J. B. Cole et al.*, AIPL, ARS-USDA

US scales

	Net merit (NM\$) (2010)	TPI (2011)
(Milk)+Fat+Protein	35	43
Longevity (PL)	22	10
Somatic cells	-10	-5
Fertility (DPR)	11	11
Calving diff./ Stillbirth	-5	-3
Type composite		10
Udder composite	7	12
Feet/legs composite	4	6
Body size	-6	




A red bracket groups the following rows: Longevity (PL), Somatic cells, Fertility (DPR), and Calving diff./ Stillbirth. The sum of their Net Merit values is 48.

A red bracket groups the following rows: Type composite, Udder composite, Feet/legs composite, and Body size. The sum of their Net Merit values is 17.

A red bracket groups the following rows: Longevity (PL), Somatic cells, Fertility (DPR), Calving diff./ Stillbirth, Type composite, Udder composite, and Feet/legs composite. The sum of their TPI values is 29.

Source 1: Net merit as a measure of lifetime profit: 2010 revision *J. B. Cole et al.*, AIPL, ARS-USDA

Source 2: http://www.holsteinusa.com/genetic_evaluations/ss_tpi_formula.html

TPI: the worst possible choice for many countries

- Very strong emphasis on type: the big “show” cow
- Not adapted to suboptimal environments !



- Most American farmers don't use it !

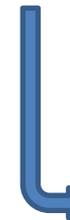
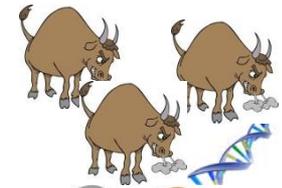
Alternatives do exist !

- Use Net Merit \$!
- Use Interbull results on a more adequate Interbull scale (New-Zealand, Ireland, South Africa, Argentina ...)
- For low input environments: choose a more sustainable Total Merit Index combining Interbull information
 - 30% Fertility, 30% Health, 20% Longevity, 10% Production, 10% Feet and legs ?
 - ➔ A much larger choice of (cheaper) bulls
 - ➔ genetic level may still be (too) high for production
 - ➔ for exporting countries: larger range of bulls

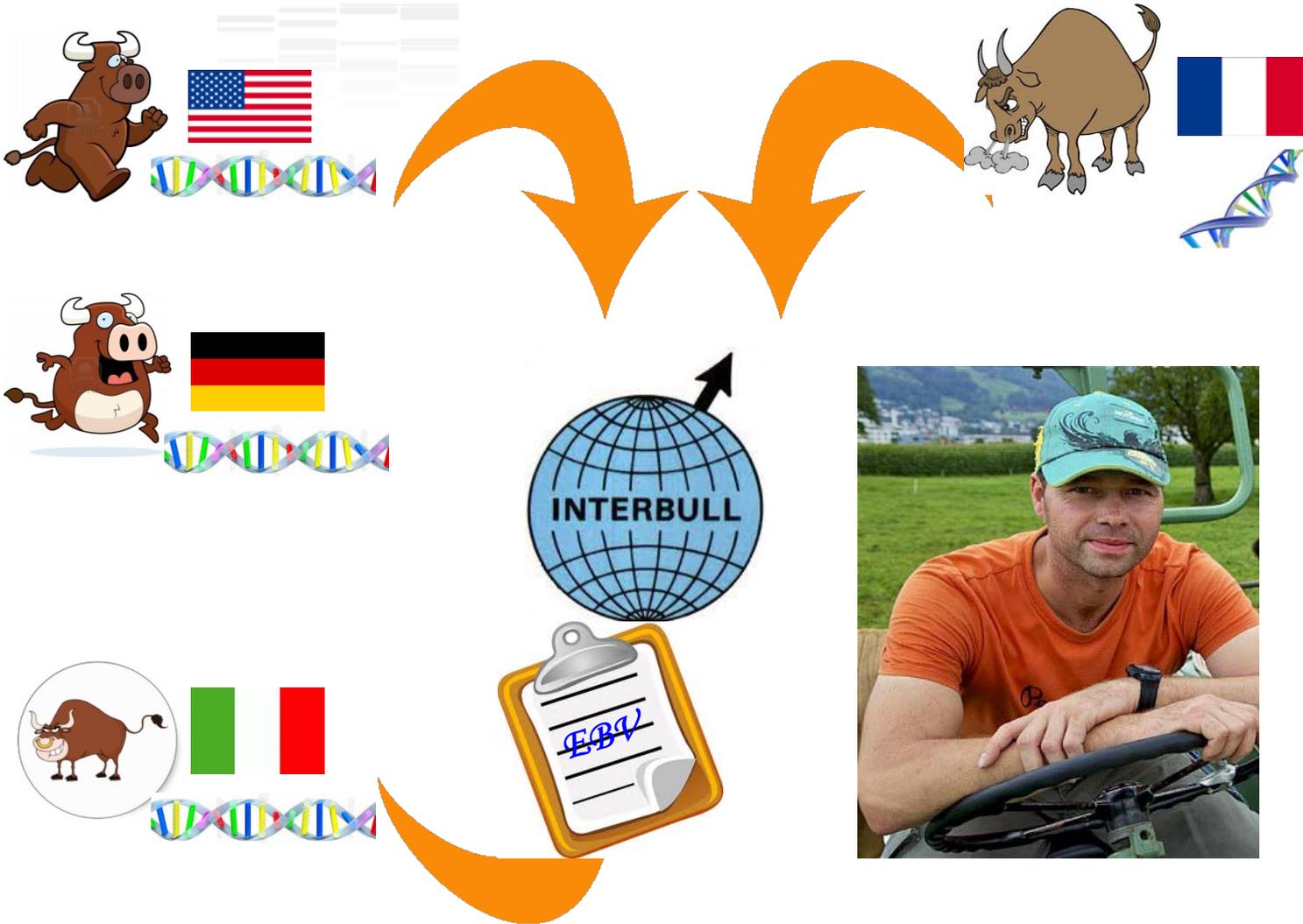
Genomic era: Interbull members



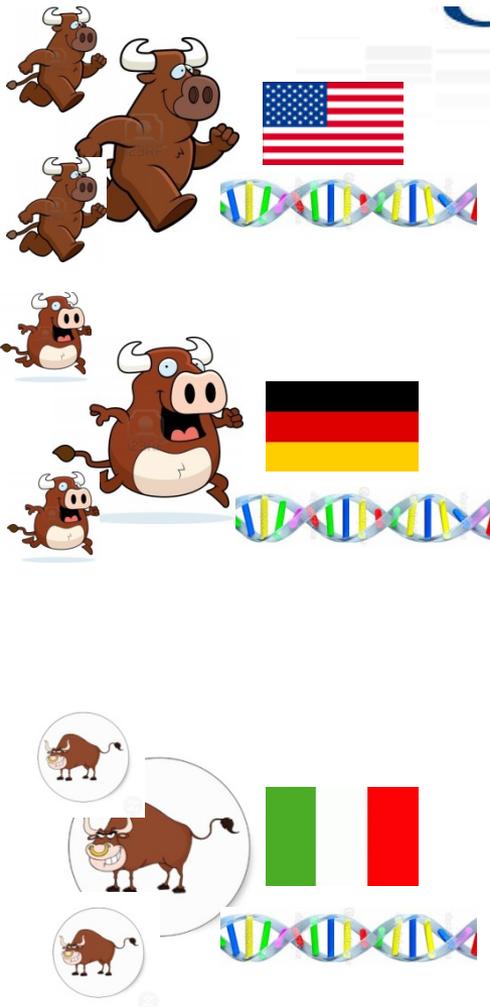
Reference population



Interbull members : the ideal situation

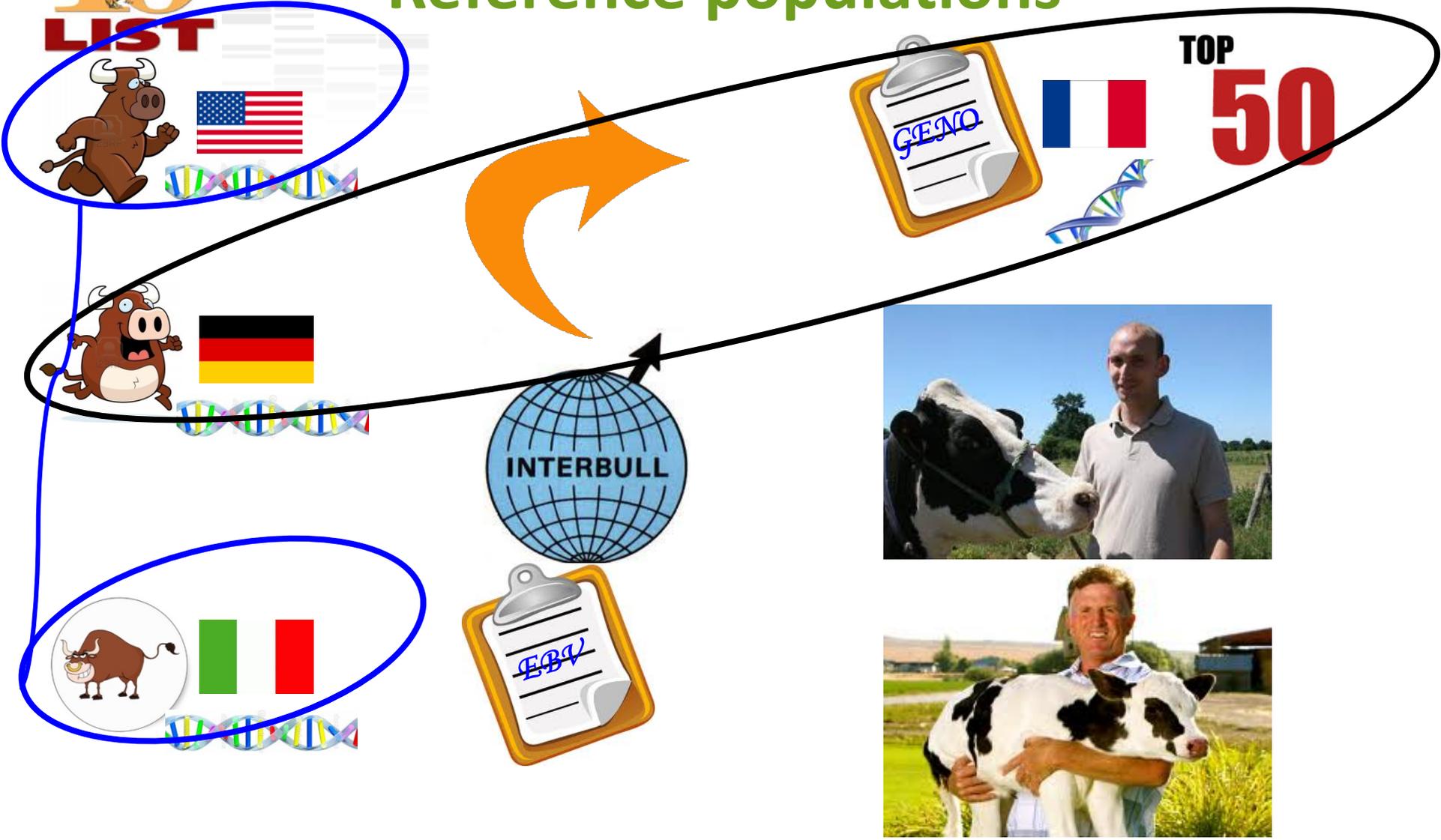


Interbull members : the ideal situation



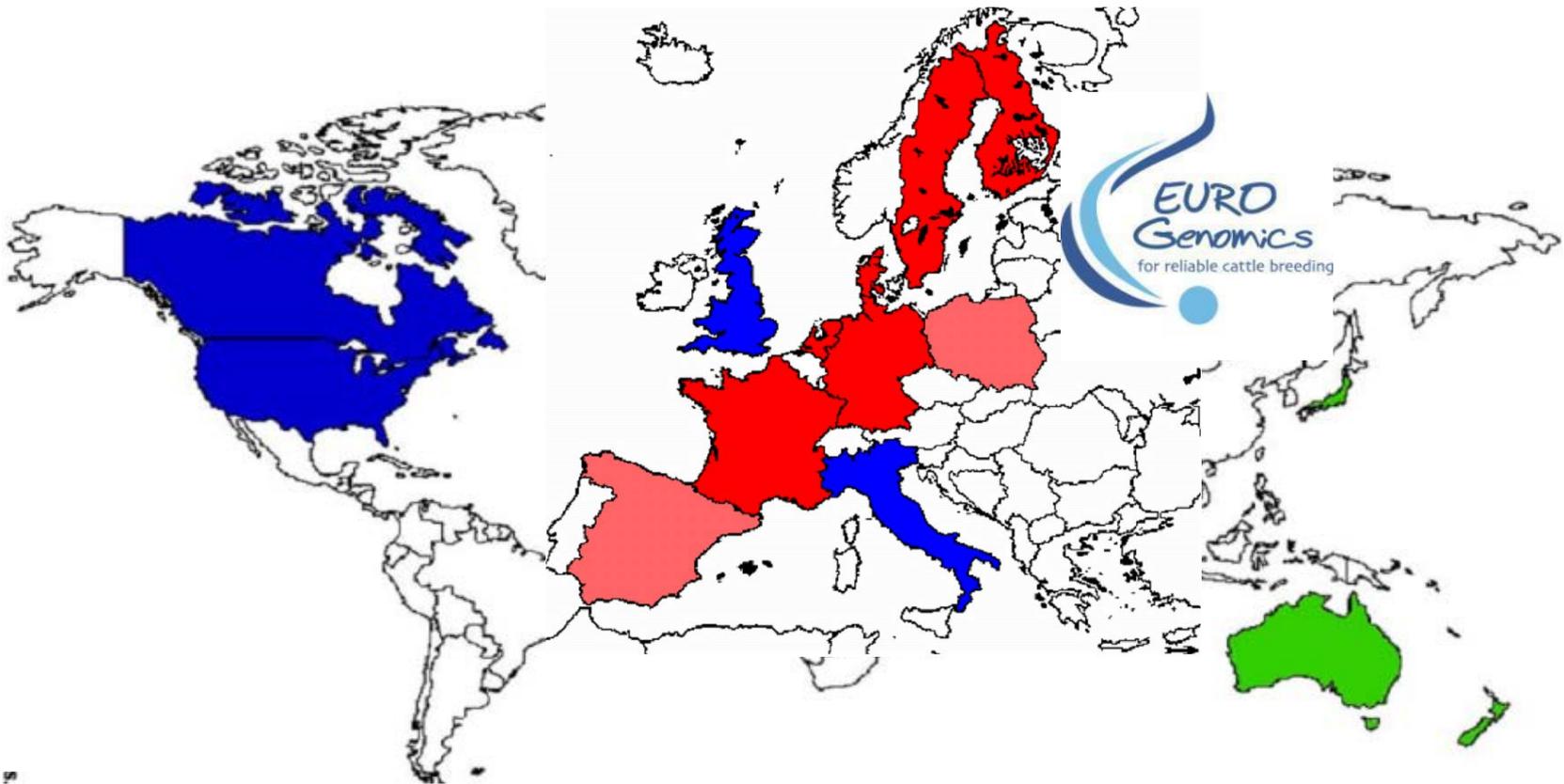
TOP 10 LIST

Reference populations



The Holstein situation

- favors the largest (groups of) countries able to jointly assemble large reference populations → 2 large consortia



Issues for countries with genomic evaluations

- Each country has its own genomic evaluation with its own particularities and its own scale
- Interbull **does not** have access to (Holstein) genotypes
- ✓ How to insure fair across country GEBV comparison?
- ✓ Interbull is developing a GMACE (extension of MACE) but faces technical + political problems

Issues for countries with genomic evaluations

- Interbull **genetic** evaluations are extremely important
 - ✓ **Only way** to combine reference populations from different countries !
 - ✓ BUT resulting **genomic** evaluations are **better** (G x E free) !

... for countries **without** genomic evaluations

- How to benefit from genomic information ?
 - ✓ How to create a national reference population large enough?
 - ✓ How to join a consortium (and which one?)
 - ✓ How to compare foreign GEBV from different countries ?
 - ✓ Rely on GMACE ? Just starting (trust?)
 - ✓ Not comparable scales → Currently the most popular is **GTPI** !



Holstein Assoc USA
@HolsteinUSA

Suivre

e.g., on tweeter:

Official Top 5 GTPI Young Bulls 5: #1
NUMERO UNO(+2604); #2
SUPERSIRE(+2581); #3
MCCUTCHEN(+2547); #4
HEADLINER(+2496); #5 MOGUL(+2493)

Répondre

Retweeter

Favori

Plus

A longer term, more **sustainable** alternative

- Help (sufficiently large) countries with performance recordings
 - ✓ **to join Interbull**
 - ✓ **and to benefit from existing reference populations**

... in order to develop genomic evaluations adapted to the local conditions and to import the best young bulls worldwide for the local needs
- e.g., INRA program GENOSOUTH with South Africa, Brazil and India

Key messages

- Genomic selection is characterized both by intense international **collaboration** and strong economic **competition**, in particular in Holstein
- Even for large exporting countries, collaborations are still needed
 - use of sequence data: 1000 genome projects
 - phenotyping of new traits (feed efficiency, methane emission)
- The gap between leading countries and the others is widening fast, to such an extent that « **collaborative spirit** » is declining



Key messages

- Fair across-country comparisons are (and will remain) a necessity
- **Interbull plays (and will play) a central role**
- For importing countries without own genetic/genomic evaluation: **a proper choice of scale is essential** and the (G)TPI is certainly NOT the most sustainable one
- For (large) importing countries, **becoming an Interbull member and developing an own genomic evaluation is strategic**
- A unique (€) scale for genetic evaluations would make Europe stronger on the international market



Acknowledgments



D. Boichard
P. Croiseau
C. Patry

S. Fritz
C. Hoze

S. Mattalia
S. Minéry
J. Promp



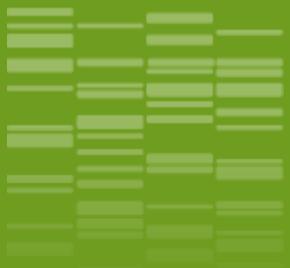
X. David

L. Journaux

LABOGENA

Thank you





International issues in dairy cattle genetics

Vincent Ducrocq

Jouy-en-Josas, France

