



Ten years of genomic selection in French dairy sheep breeds: assessment and perspectives

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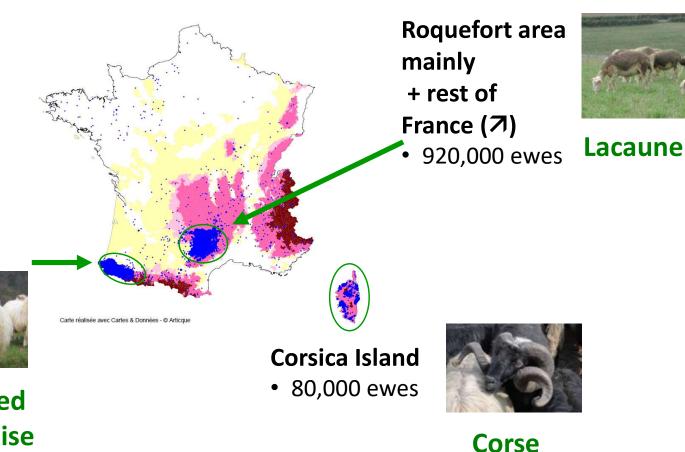






Dairy sheep populations in France 1,450 M dairy sheep in France (30% of sheep population)

Local breeds raised in their own areas (mostly harsh & mountainous) & production systems.



Western Pyrenees

• 450,000 ewes







Black-faced & Red-faced Manech, Basco-Béarnaise

The selection programs in French dairy sheep breeds: main features and current situation

 Pyramidal organization within each breed for benefit to the whole population (organize both creation and diffusion).

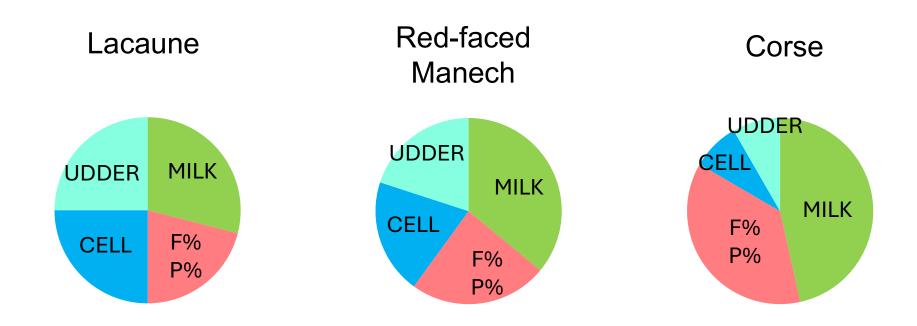
Lac	caune	Basco- Béarn.	BF Manech	RF Manech	Corsica
# flocks in selection program	205 k	29 k	9 k	79 k	16 k

Major role of Al

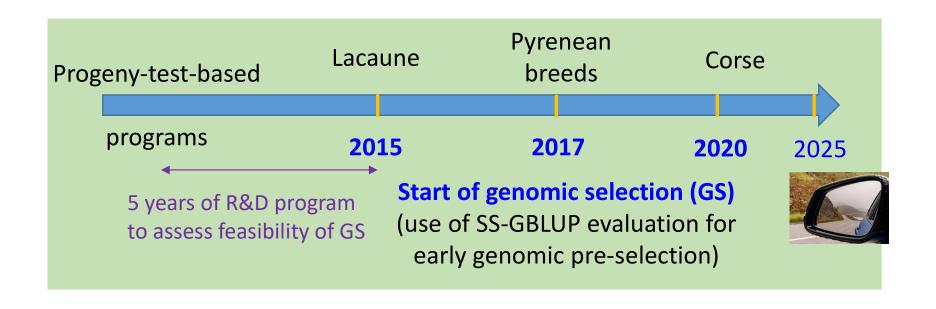
% AI in selection flocks	84%	47%	43%	47%	46%
# new AI rams each year	320	70	25	160	20
# total AI rams in AI centre	1100	180	75	420	45
# young rams genotyped	3700	300	85	800	350

Balanced breeding goals

- The selection objectives are more and more balanced
- => Selection criteria include a growing variety of abilities / traits.
- Weights adapted to each breed

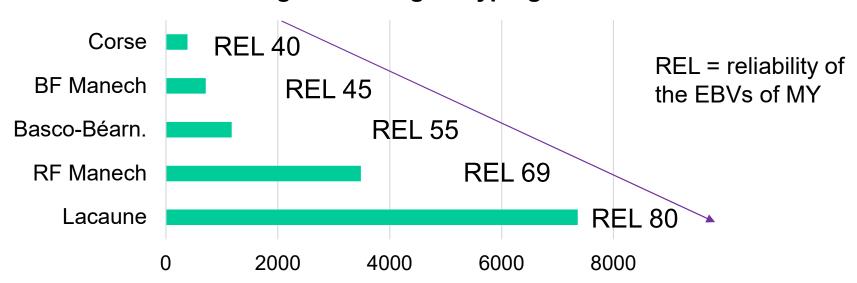


Timeline of genomic selection in French dairy sheep



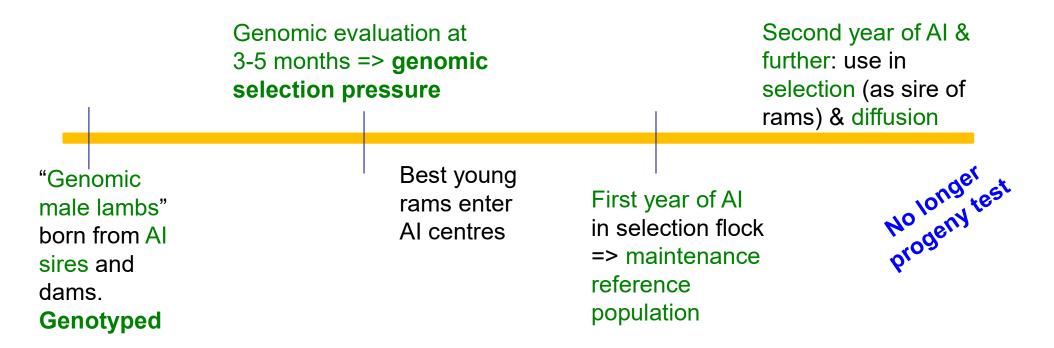
Size of the reference populations / consequences on reliability

Rams with daughters and genotypings



Reliability significantly increases with size of reference population

How does genomic programs work?



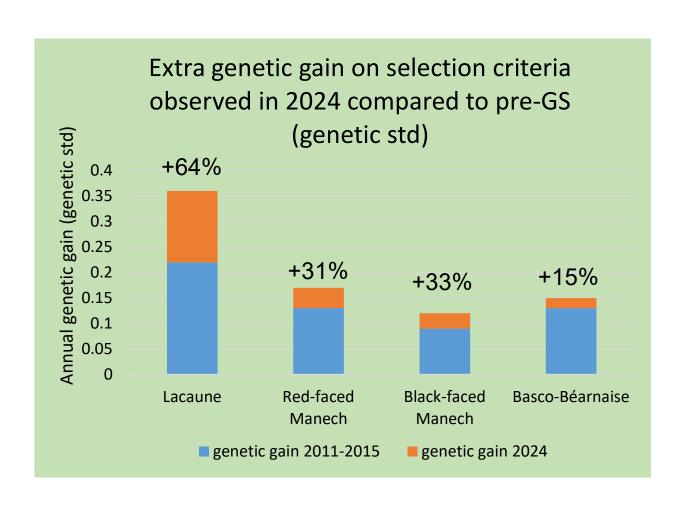
Actual genomic selection pressure << apparent (# rams selected / # candidates) Lacaune ~ 20%

BB ~ 58% BFM ~ 48% RFM ~ 53%

Corse ~ 52%

Benefits of genomic selection

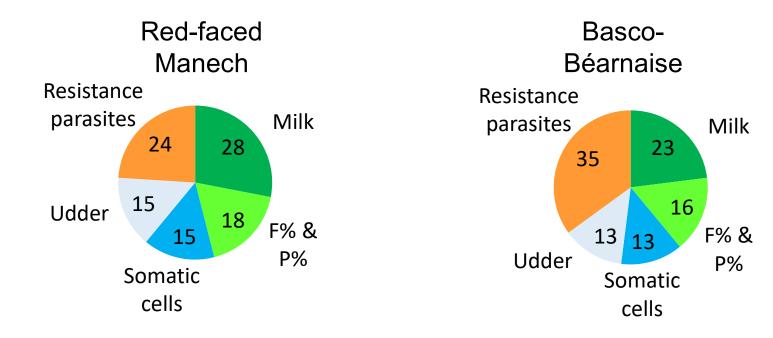
- Generating an annual genetic gain for economical index ranging from 0.12 to 0.36 genetic standard deviation
- Increase in genetic gain from 15 to 64%



How to valorise a most efficient genomic selection?

Extra genetic gains permitted by GS is / will be oriented toward inclusion of new traits related to adaptation / resilience / efficiency

Since 2024, resistance to parasites included in the selection objective of Basco-Béarnaise and Red-faced Manech



- Other actual selected criteria: horn, functional morphology (including feet and legs), PrP, semen production
- On-going criteria: longevity
- R&D criteria: feed efficiency, GHG emissions, fine milk components (caseins)

Multi-purpose valorisation of genotypings

Besides GS:

- Parentage verification followed by parentage discovery applied on males chosen for genomic selection (before genomic pre-selection)
 - √ 4-5% of wrong sires => ~90% of sire discovery
- **Sire discovery** on ewes in flocks that does not do AI (including organic farms where synchronisation by hormone is forbidden)
 - \checkmark 3,500 ewes in 2023 => ~95% of sire assignation

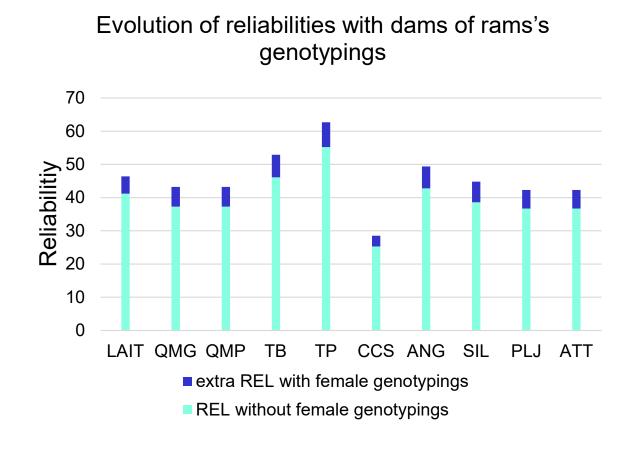
Major genes

- ✓ PRP scrapie resistance
- ✓ SOCS2 susceptibility to somatic cells
- ✓ Horn management of horn in Red-Faced Manech (& Corse breed)
- ✓ New genes in the next years lethal mutations, cryptorchidism
- Better management of inbreeding
 - ✓ Not yet in routine in France

Genotypings of females: complement the reference population

Case of Corsica breed

- Corsica breed: few rams in the reference population – 385 rams, +20 each year
- Decision to switch toward a balanced selection objective
 => Phenotyping of milk quality / cells / udder started in 2017
- Very low reference population on new traits
- Genotypings of dams of rams allow an increase in precision of EBVs

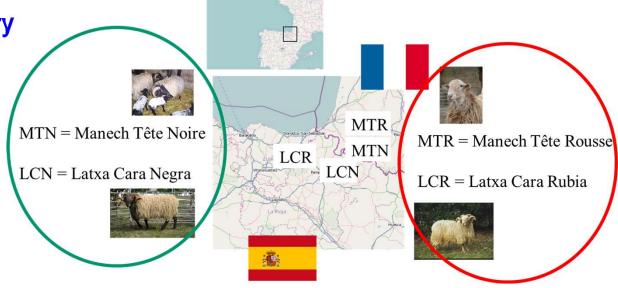


~2,100 dams of rams genotyped Reliability: gain of 5-7 points

Toward a genomic selection at a multi-country level

Collaboration Basque Country

Spain / France



A series of projects since 2010 : Genomia, ARDI, SMARTER, ARDI2

- Creation of ARTALDEAN ~ across country breed society
- Across-country genomic evaluation
- Genotypings merged/processed in GenoEx in Interbull







Sheep GenoEx (ARDI2 specifically financed the project in Interbull for its needs) might be expanded to other situations in small ruminants.

Take home messages

- ✓ Genomic selection started in France in 2015 in Lacaune and spread quickly to the other breeds.
- ✓ After 10 years, an extra genetic gain is observed in mostly all breeds. Highly fostered by organised and collective breeding programs.
- ✓ Genomic selection cost-effective: higher genetic gain, decreasing cost of genotypings compared to 10 years ago, existing market for "genomic" rams destined to natural mating, multi-purpose valorisation of genotypings.
- ✓ Higher efficiency used to include new traits in the selection program (e.g. resistance to parasites, functional morphology)

Pending questions and perspectives

- ✓ Use of genomic selection at an across-country level: increase reference population, compensate smaller populations in selection than in cattle, increase accuracy and selection intensity
- ✓ **Local** breeds vs "international" breeds
- √ Various issues: technical, legal (sharing of data)
- Existing initiatives to promote and facilitate networking and "actions" across countries
- ✓ ICAR sheep, goat, camelid WG proposed an EU Reference Centre in small ruminants (with outcomes beyond EU) to produce harmonisation, services and incentives in the field of performance testing and genetic evaluation in sheep and goats, with focus on genomics

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