

## WP3: Genetics of trade-offs and synergies between resilience and efficiency related traits

Major achievements

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#### **Objectives of the WP**

The challenge for livestock breeding is to improve the resilience traits simultaneous with feed efficiency and other traits important for a sustainable livestock sector. Simultaneous breeding for multiple traits can fall foul of trade-offs between traits. The WP objectives are to:

- To identify trade-offs & synergies under genetic control (Task3.1)
- To better understand the biological mechanisms underlying those trade-offs & synergies and how they affect resilience and efficiency (Task3.2)
- To develop prediction models to manage trade-offs and optimize resilience and efficiency in challenging conditions (Task3.3)



#### Task 3.1. Identify trade-offs under genetic control

- Systematic review on host genetic component traits underlying herd disease resilience, and the influence of trade-offs on it (UEDIN) :
  - Published here : Doeschl-Wilson et al. "Livestock disease resilience: From individual to herd level" Animal, Dec Vol. 15. Suppl 1, 2021
  - Potential trade-offs between individual resilience and herd resilience
- □ Meta-analysis of genetic parameters (SRUC, INRAE, UEDIN)
  - 2,151 estimates of heritability and genetic correlation for 81 sheep and 14 goat breeds collated from 12 SMARTER partners AGRIS and public data.
  - Results were published here : Mucha et al. « Animal Board Invited Review: Meta-analysis of genetic parameters for resilience » Animal, Vol16, issue3, 2022
- Genetic analyses for pleiotropic QTLs and potential underlying genes and pathways associated with efficiency, resilience and reproduction related traits (INRAE, UNILEON, INIA\_UY)

Smarter

Limited evidence of genetic antagonisms found between resilience and efficiency for dairy goats (SCS with fat content) and dairy sheep (SCS with protein content), and not for meat sheep. => Selection for both R&E is feasible. Even though the pooled estimates were non significant, antagonisms may exist but only in specific populations and environments (Figures a & b)

Forest plots showing genetic correlation estimates between:









Mucha et al. (2021) Animal

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## Task 3.1. Identify trade-offs under genetic control

- pleiotropic QTLs and potential underlying genes and pathways associated with efficiency, resilience and reproduction related traits
- 7 GWAS datasets collated (or newly generated) (UNILEON, INIA-UY, Capgenes, INRAE) from 8 breeds (Romane, Lacaune, Saanen, Alpine, Churra, Corriedale, Merino, Texel) with > 15000 genotypes
- Pleiotropy identified by Clip Test (David et al., 2023)



- GWAS Pleiotropic QTLs identified (CNR, INRAE) :
  - ✓ Lacaune : CHI3 for mastitis- milk production
  - ✓ Saanen: CHI19 for mastitis-longevity-production
  - Romane: OAR1 for Body Condition Score and Body weight

Source: Johanna Ramirez-Diaz (CNER), 2023



□ Five data sets were created in genetically selected sheep and goat which will undergo both nutritional and infection challenges

	Breed Species (Partner)	Genetic group	Challenge	Status
And a	ASSAF <b>Dairy sheep</b> (UNILEON) N=60	Milk production	Nutritional : growing period (N=60) Infectious: LPS* in L1 (N=24)	Complete
	ALPINE <b>Dairy goat</b> (INRAE) N= 98	Longevity	Nutritional: 2 days on hay in L1 (N=98) Infectious = LPS* (N=87)	Complete
A AT	ROMANE <b>Meat sheep</b> (INRAE) N=48	Parasite resistance	Infectious: parasite <i>H. contortus</i> Nutritional: low/high protein around lambing	Complete
(A)	ROMANE <b>Meat sheep</b> (INRAE) N=60	Feed efficiency	Infectious: parasite <i>H. contortus</i>	Complete
ANA	CORRIEDALE <b>Meat sheep</b> (INIA-UY) N = 67	Parasite resistance	Nutritional: None (feed intake records) Infectious: parasite <i>H. contortus</i>	Complete

#### \*LPS: challenge with lipopolysaccharide mimicking an inflammatory disease such as mastitis

□ Results in **ROMANE meat sheep (INRAE)** genetically selected for parasite resistance

Selection for resistance to gastro-intestinal parasite in growing lambs effective

- most of the time in adult breeding ewes  $\checkmark$
- at peripartum X

Around lambing, level of protein in the diet affects

- ewe growth  $\checkmark$  (and lamb growth  $\checkmark$ )
- ewe resistance to parasite X





Trade-offs between resistance to parasite and production

- during growth ✓ (reduced fat deposition) <</li>
- during reproduction X (except around 1<sup>st</sup> lambing where a cost of host genetic resistance on body weight has been detected)



□ Results in **ROMANE meat sheep (INRAE)** genetically selected for feed efficiency



No line difference in FEC during growth No effect of infectious challenge on residual feed intake (RFI)

Trade-off between feed efficiency and parasite resistance

- during growth X
- during reproduction ? (data analysis in progress)

Douhard et al. "Feed efficiency and resource allocation trade-offs: theory, evidence and prospects" Proceedings of 12<sup>th</sup> WCGALP



□ Results in ASSAF Dairy sheep (UNILEON) in genetically selected for milk production



Evidence for an effect of DIET and genetic LINE on the outcome of the inflammatory challenge with LPS (Cytokine/chemokine ratio traits) : DIET significant for several plasma markers: IL-10, IL36RA and VEGF-A. LINE significant for a number of markers : IL-8.

=> Analysis of mechanisms that underlie the trade-offs are in progress





#### https://doi.org/10.1016/j.rvsc.2023.04.006

Smarter

Influence of a temporary restriction of dietary protein in prepubertal ewe lambs on first lactation milk traits and response to a mammary gland inflammatory challenge

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#### Influence of dietary protein restriction in prepubertal ewe lambs on first lactation milk traits and response to a mammary gland inflammatory challenge



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as a potential cause mediating these results.





1. RNA-Seq of Milk somatic cells (MSC) of Control animals (no protein restriction in prepubertal ages ) Dynamic milk transcriptome expression changes in response to *E. coli* LPS intramammary challenge: Differentially expressed genes (DEGs): **13 NC vs 11 C** 



Three groups of DEGs were defined:

- Acute inflammatory response DEGs (A+C): 521 genes with higher expression at 6h
- *General inflammatory response DEGs* (B): 1436 genes with higher expression at 6h and 24h
- Late inflammatory response DEGs (D): 67 genes with higher expression at 24h





□ Results in **CORRIEDALE sheep (INIA-UY)** genetically selected for parasite resistance



Analysis of feed conversion efficiency in Corriedale lines (Ferreira et al. 2021)

Trait	GIN line				
	Resistant	Susceptible	Р		
RFI	$0.02\pm0.018$	$-0.02 \pm 0.016$	0.116		
FCR	$9.0\pm0.62$	$7.6\pm0.75$	0.161		
DMI	$0.97\pm0.036$	$0.98\pm0.044$	0.969		
ADG	$123\pm0.90$	$143\pm0.11$	0.168		
Initial BW	$27.1 \pm 0.53$	$27.7\pm0.64$	0.483		
Final BW	$33.9\pm0.46$	$34.7\pm0.54$	0.292		

Corriedale lines did not show significant differences in residual feed intake (RFI), feed conversion ratio (FCR), dry matter intake (DMI), average daily gain (ADG) or body weight (BW), indicating that breeding parasite resistant animals in that population would not have undesirable effect on feed efficiency.



# Task 3.3: Develop prediction models to manage trade-offs and optimise resilience & efficiency in challenging conditions

**Objective of the task:** Develop mechanistic and statistical models to predict and manage trade-offs associated with infectious and nutritional challenges

- **1. Marie Ithurbide** (INRAE, Toulouse): Toward genetic selection for resilience based on milk metabolites
- 2. Nicolas Friggens (INRAE, Mosar): Towards a resource allocation model for dairy goats building on SMARTER data
- **3.** Frederic Douhard (INRAE, Toulouse): Modelling the energy cost of host resistance to gastrointestinal parasites in meat sheep
- 4. Masoud Ghaderi-Zefreh (Roslin Institute): Evaluation of novel resilience phenotypes based on longitudinal performance measures



## Thank you !