

➤ How genetic selection could contribute to the multi performance of farming systems?

A modelling approach applied to Lacaune dairy sheep

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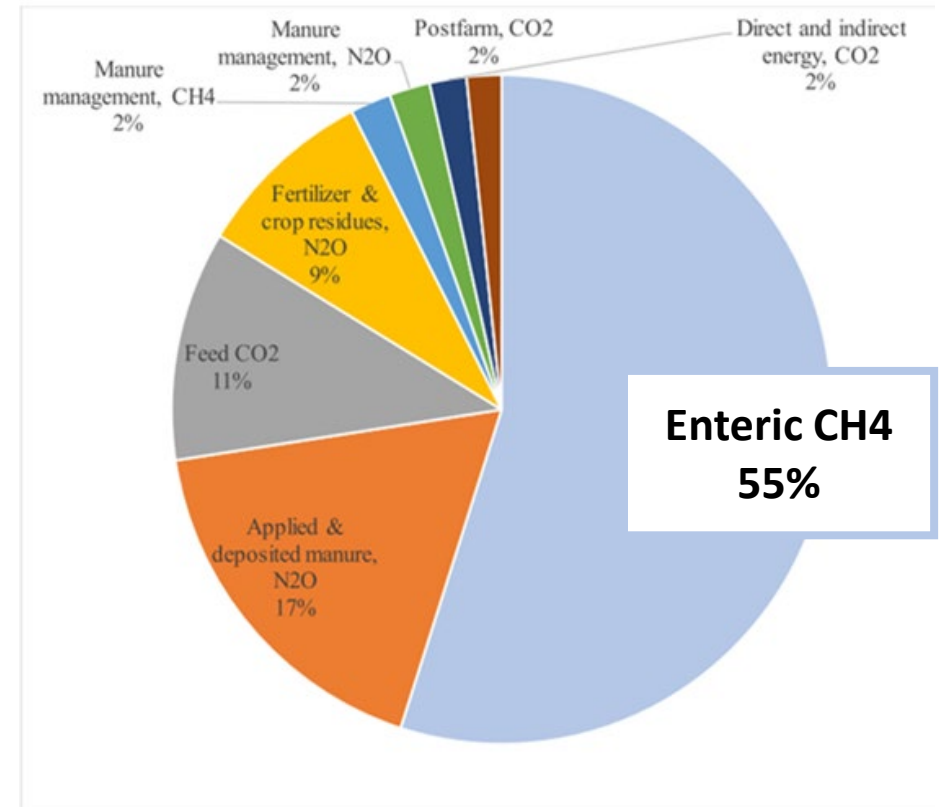
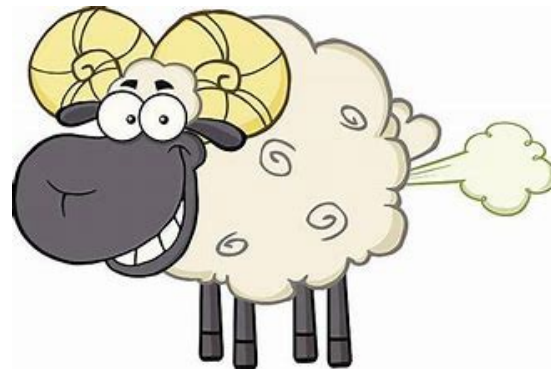
Multiperformance of farming systems

Sustainability (Clay et al. 2020)

Economic
efficiency

Social impact

Environmental
impact



Global GHG emissions from small ruminant supply chains; adapted from Gerber et al. (2013).

Improvement approaches

Multiperformance

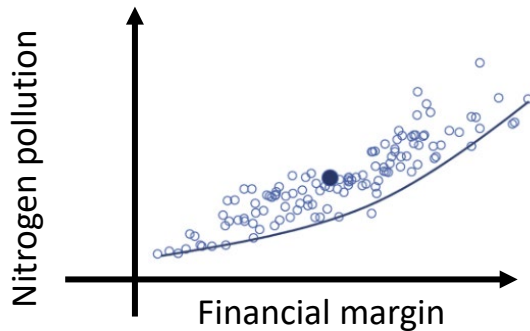
Management practices Farming system design

- Feed, reproduction

Selection practices

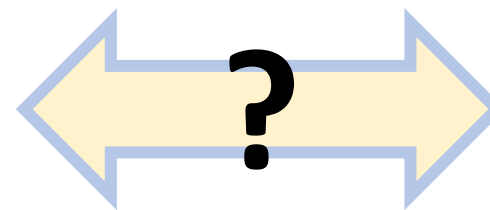
- Selecting traits that enhance both productivity and sustainability
 - Operates at **population** levels
 - Focusing on **long-term** changes

→ use of **individual-based models** to manage trade-offs at farm level



Ex : in dairy sheep
Villalba et al. 2019

Lacaune breed



Current
breeding
goal

Milk yield

Milk composition

Udder score

Somatic cell score

→ no selection ongoing for environmental performances (e.g. CH4)

Approach of this study

Individual-based modelling to predict responses to variation in production environments and to selection
→ trade-offs at the individual level based on resource allocation theory

Examples:

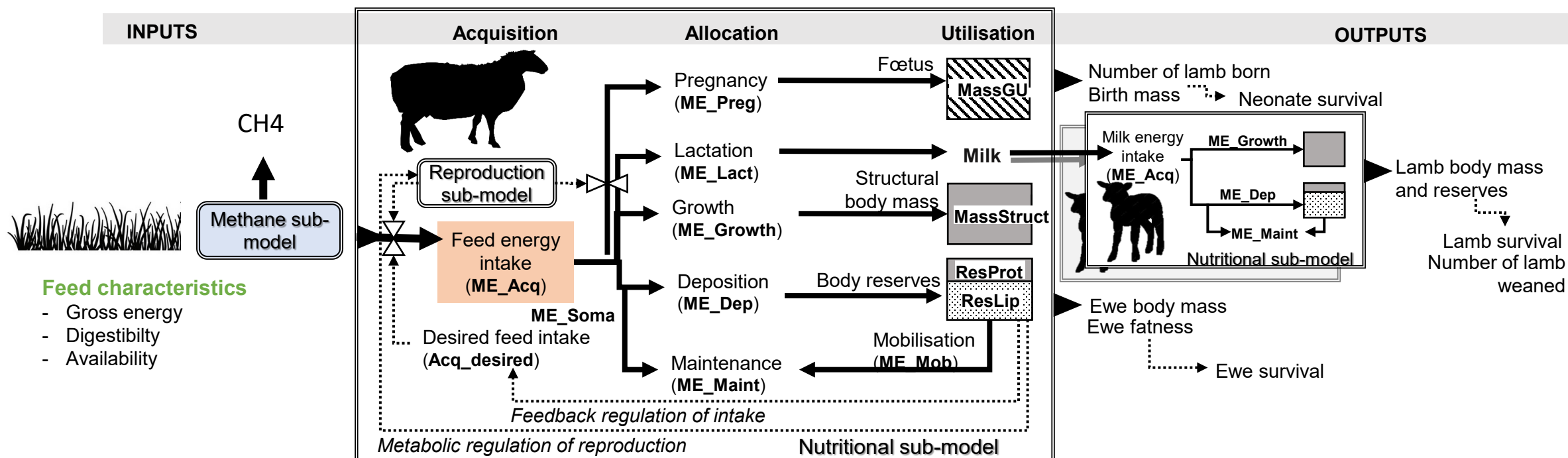
Dairy goats: *Douhard et al. 2014*

Dairy cows: *Puillet et al. 2016, 2021; Bouquet et al. 2024*

Aim: Simulate phenotypic variability of the output traits using an individual-based model of dairy sheep



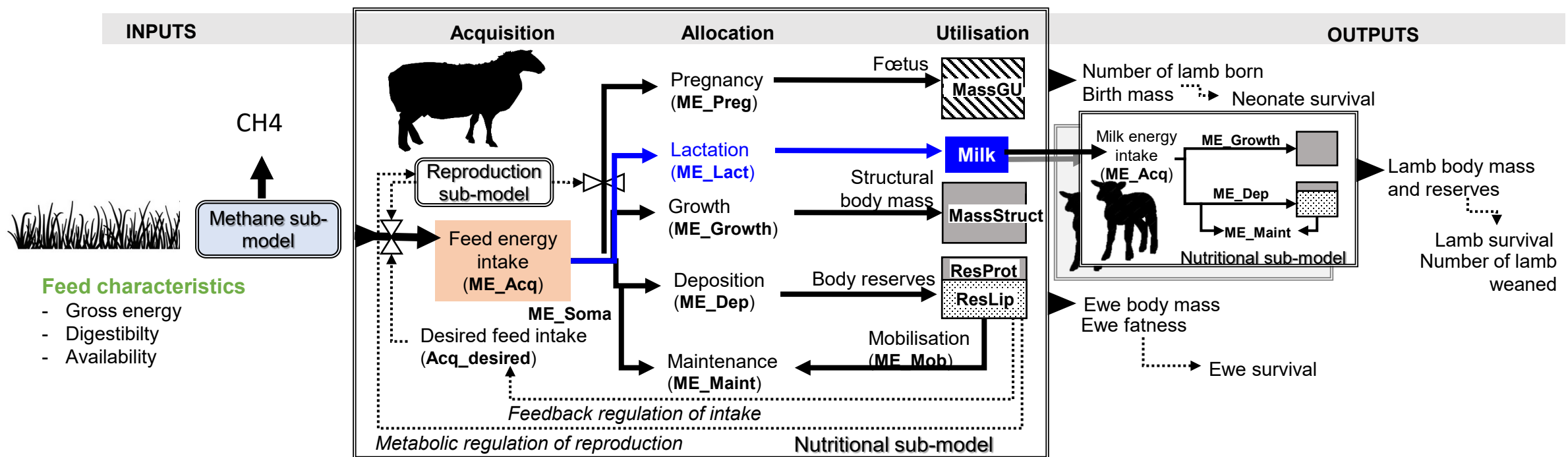
Sheep model



Metabolizable energy intake (ME_{Acq})

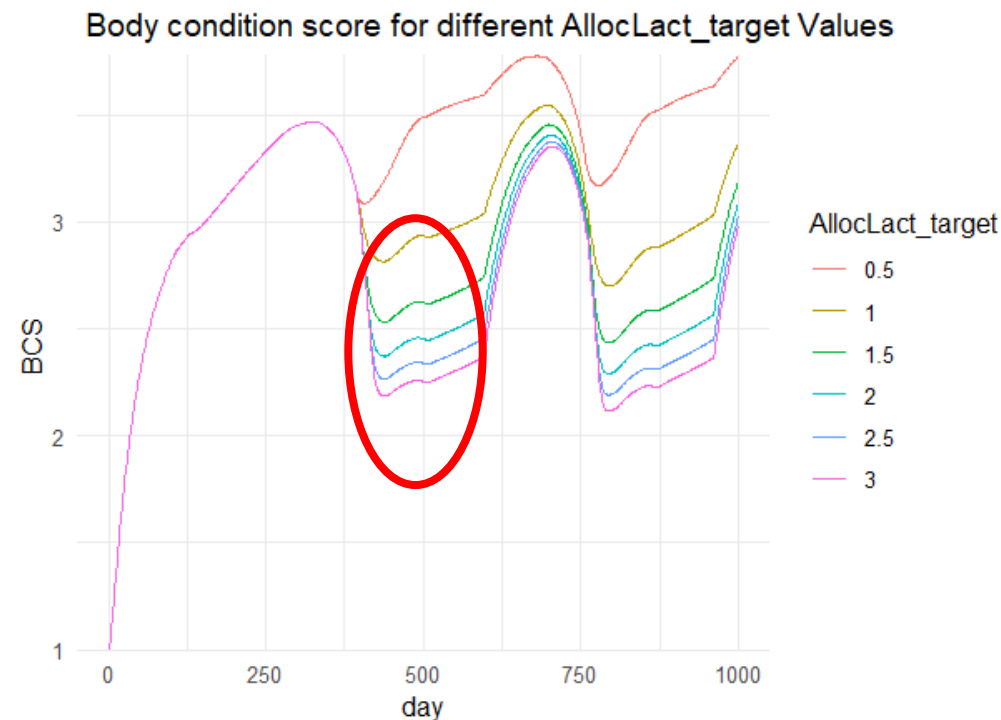
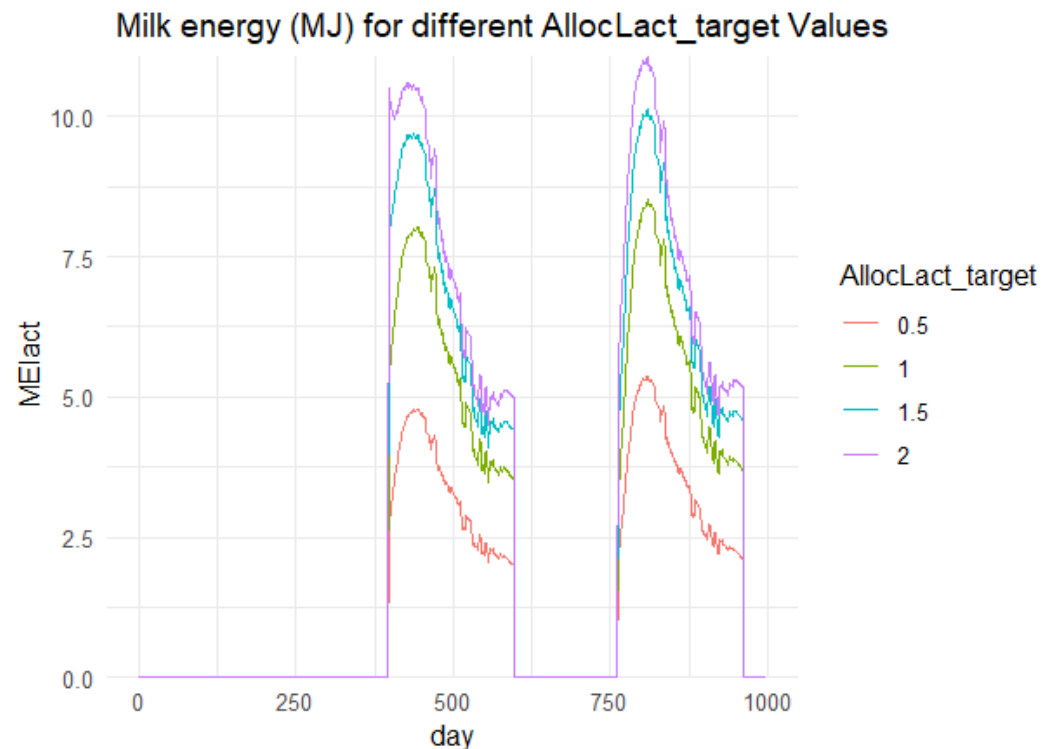
= Feed gross energy – Gaseous energy (CH₄) – Urinary energy – Fecal energy

Focus 1: simulation of individual variability



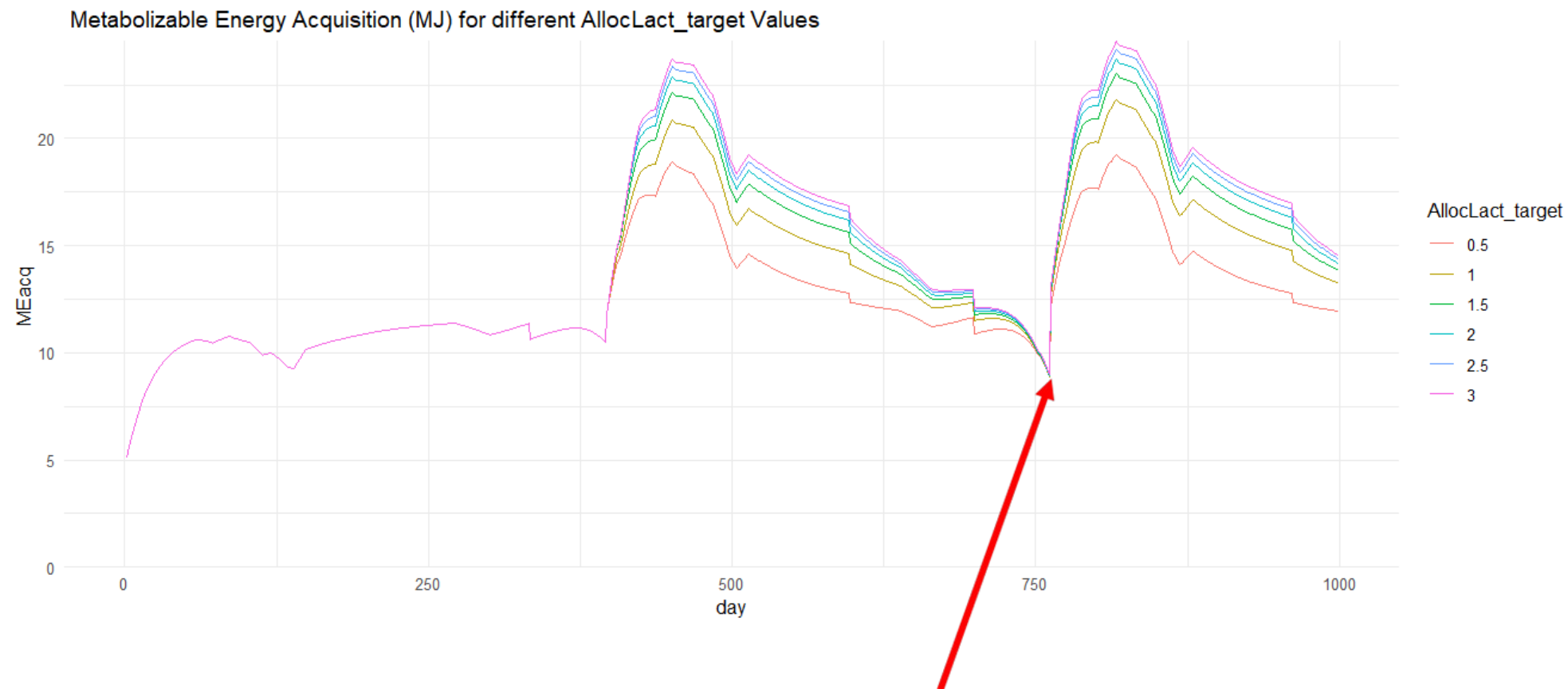
example : $ME_{Lact} = ME_{Acq} * AllocLact$
 with $AllocLact = AllocLact_{target} * f(Lact_time)$

Focus 1: simulation of individual variability



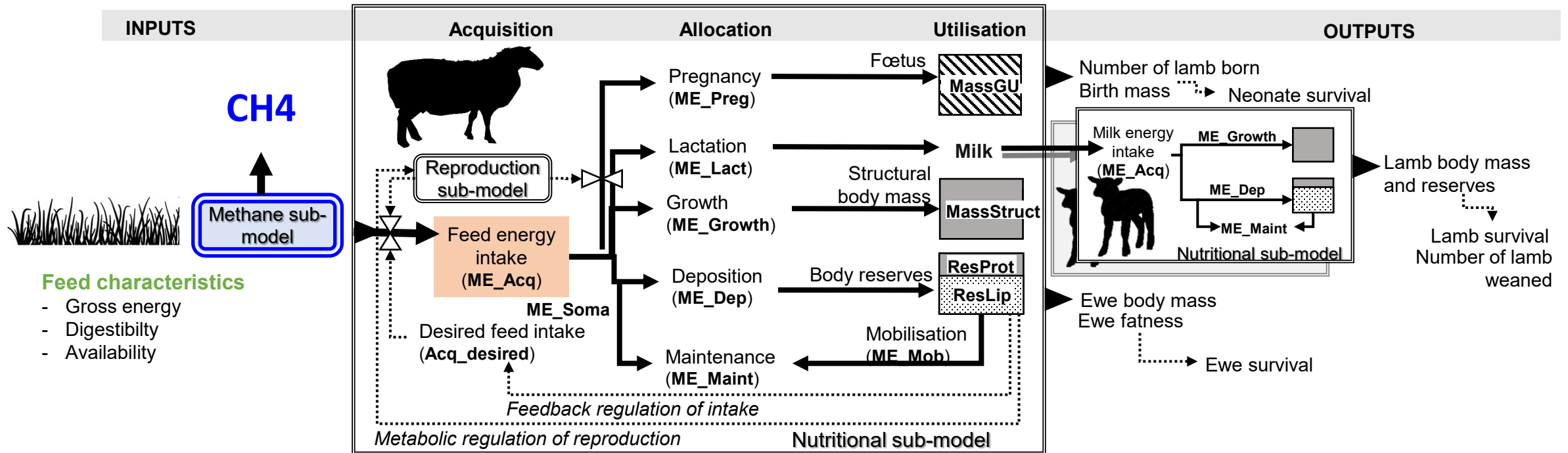
→ **AllocLact_{target}** can mediate a trade off among individuals between milk and BCS

Focus 1: simulation of individual variability



Reduction of ewe intake at the end of gestation when the gravid uterus mass is high (*Wilkinson and Chestnutt 1988*)

Focus 2: predict enteric CH₄ emissions



Metabolizable energy intake (ME_Acq)

= Feed gross energy – Gaseous energy (CH₄) – Urinary energy – Fecal energy

Focus 2: Predict enteric CH₄ emissions

$$\text{CH}_4 = \text{DOM} \times \left(45.42 - 6.66 \cdot \left(\frac{\text{Feed Intake}}{\text{Bodyweight}} \right) + 0.75 \left(\frac{\text{Feed Intake}}{\text{Bodyweight}} \right)^2 + \right.$$

Feed characteristics:

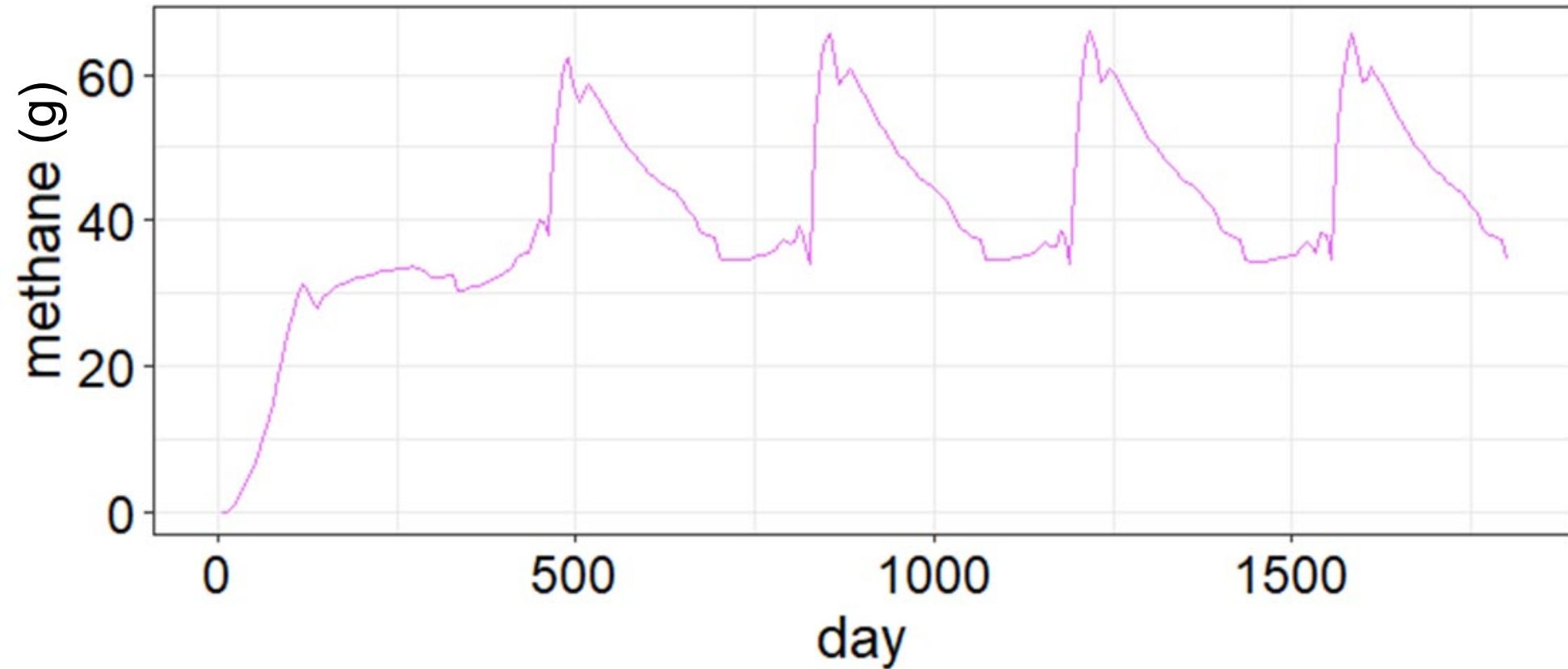
- Digestible organic matter
- Proportion of concentrate in feed

$$19.65 \cdot \text{PCO} - 35.0 \cdot \text{PCO}^2 - 2.69 \cdot \left(\frac{\text{Feed Intake}}{\text{Bodyweight}} \right) \cdot \text{PCO})$$

Sauvant et al. 2011

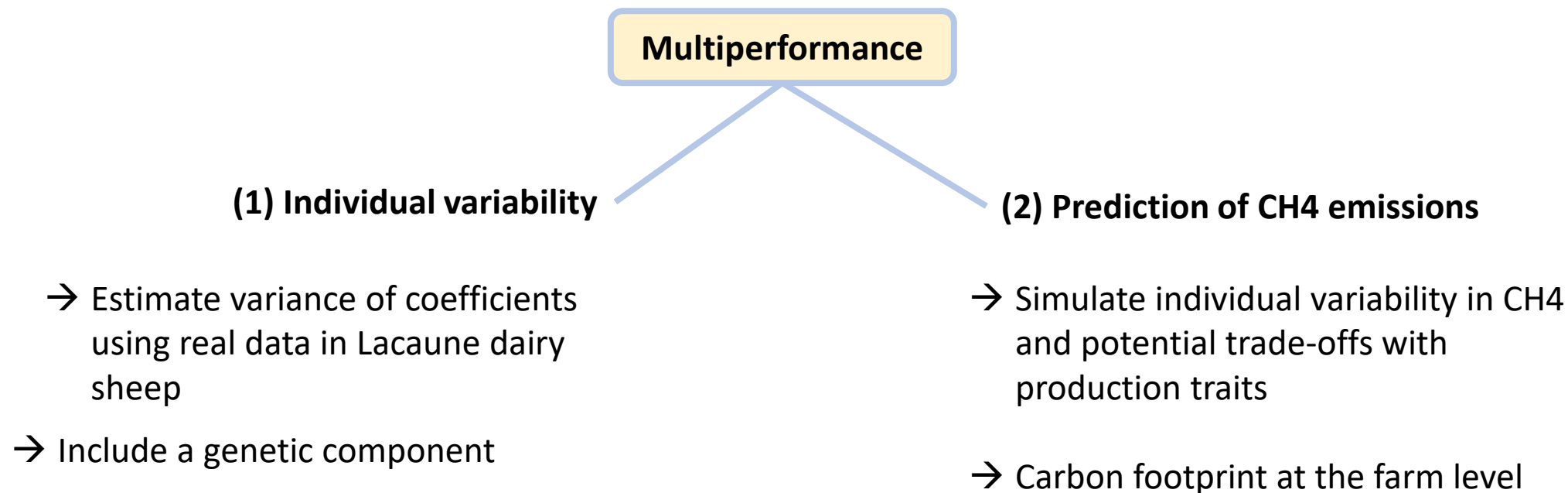


Focus 2: Predict enteric CH₄ emissions



→ Within-individual changes in enteric CH₄ emissions can be predicted

Conclusions & the next steps



Example : h^2 **AllocLact**_{target}

Bouquet et al. 2024

→ An approach to explore how genetic selection could contribute to the multi performance of farming systems



