

NATURAL ^{15}N ABUNDANCE OF ANIMAL PROTEINS: A PROMISING BIOMARKER OF FEED EFFICIENCY IN BEEF CATTLE



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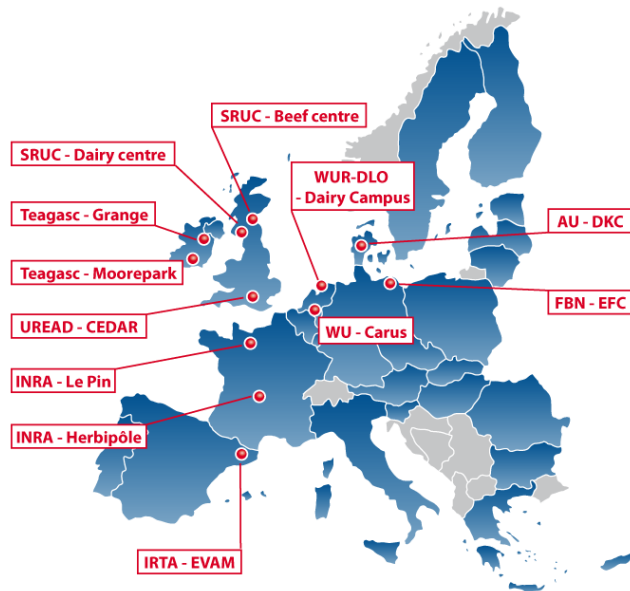
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INRAE



AN INTEGRATED INFRASTRUCTURE FOR INCREASED RESEARCH CAPABILITY AND INNOVATION IN THE EU CATTLE SECTOR



Key EU cattle research infrastructures:

- 7 countries
- 11 research infrastructures (18 installations)
- 3500 cattle

Work package 6:

Developing and evaluating promising biomarkers to predict feed efficiency and its determinants

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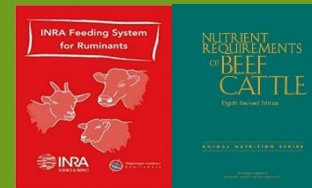
The SmartCow project has received funding from the European Union's Horizon 2020 research and innovation programme under grant agreement N°730924



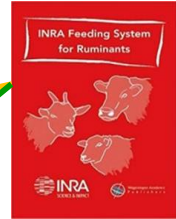
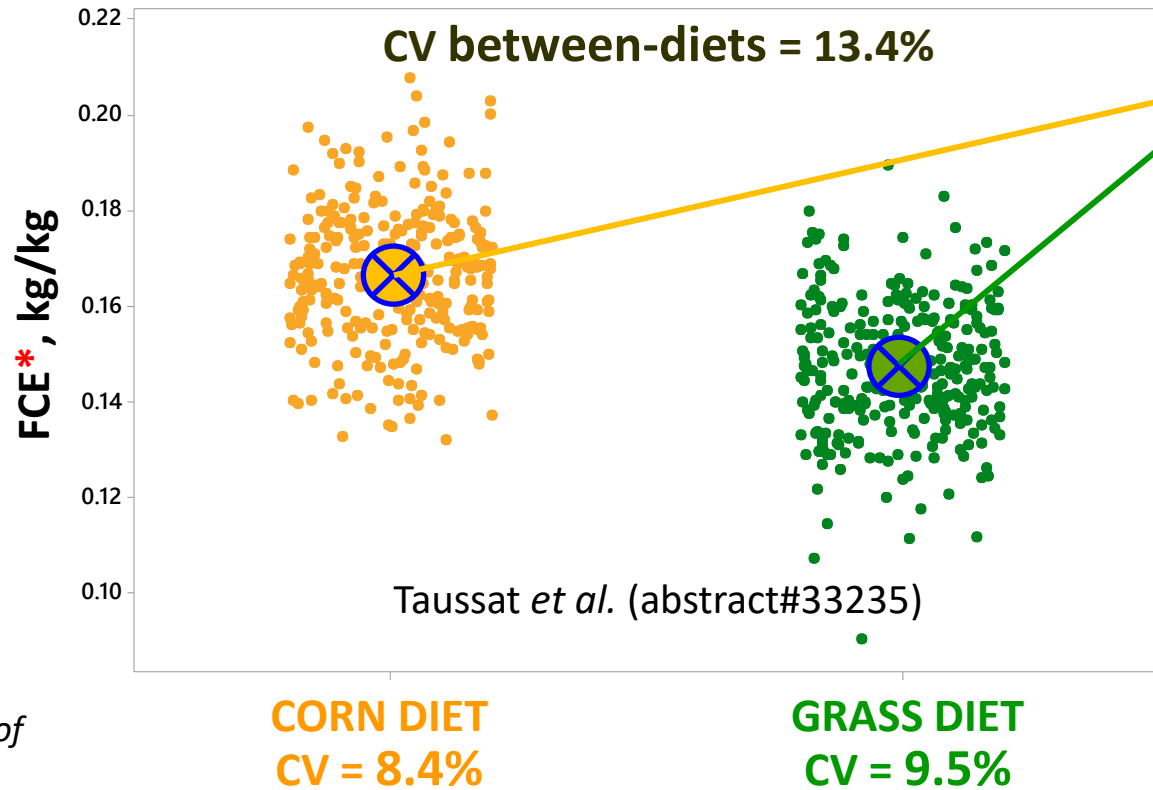
CONTEXT (i)

Improving animal feed efficiency (FE) may contribute to the sustainability of the beef cattle sector (Hill, 2012)

- NUTRITION: « DIET as a driver » ➡ Group of animals
- GENETICS: « ANIMAL as a driver » ➡ Between-animal variation
- NUTRITION × GENETICS: « One ANIMAL, one DIET » ➡ Precision feeding



NUTRITION VS GENETICS



*adjusted for the effects of
the contemporary group



CONTEXT (ii)

Phenotyping FE is costly, labor- and time-consuming !!
.....and sometimes not feasible (grazing)

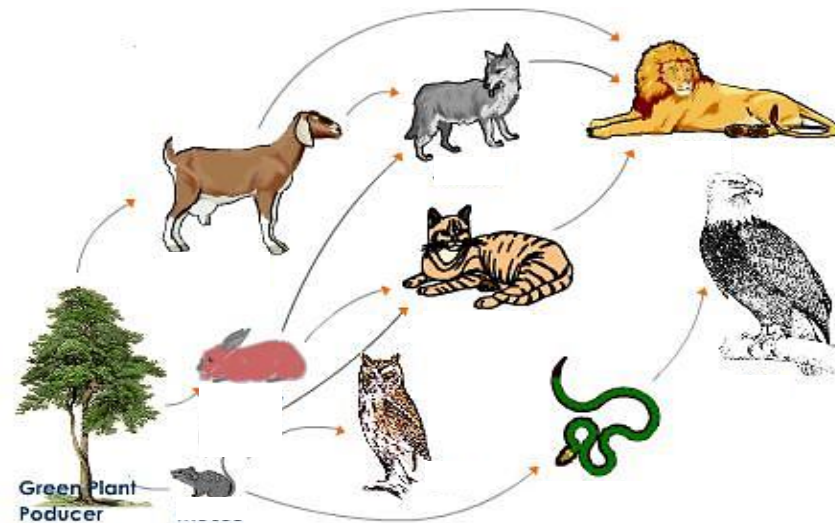
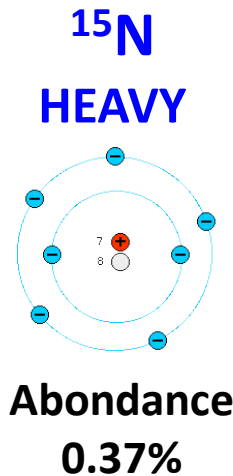
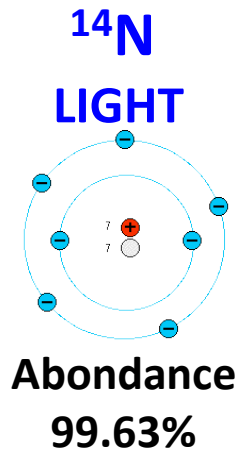
- => PROXIES to assist the genetic selection and precision feeding
- => None have been proposed & validated for predicting FE in beef cattle (yet genomics appears promising)

ISOTOPIC NITROGEN DISCRIMINATION has a potential to reflect animal-to-animal variation in FE in beef cattle
(*Wheadon et al., 2014; Cantalapiedra-Hijar et al., 2015*)

WHAT THE H*LL IS ISOTOPIC DISCRIMINATION?



NITROGEN ISOTOPES IN THE NATURE



DIFFERENT ISOTOPE DISTRIBUTION

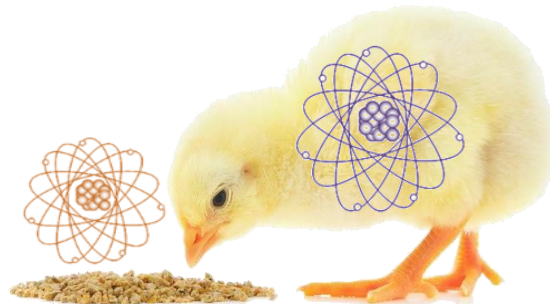
ISOTOPIC TRANSFER FROM DIET TO TISSUES

WE ARE WHAT WE EAT



NOT COMPLETELY TRUE FROM
AN ISOTOPIC POINT OF VIEW!

BUT



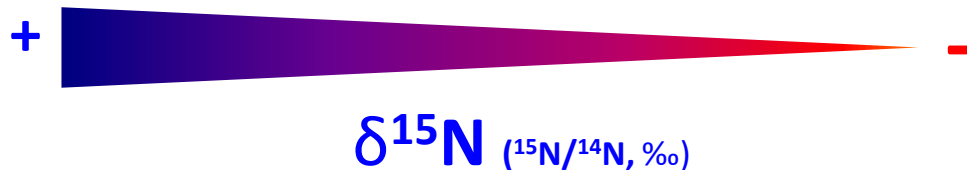
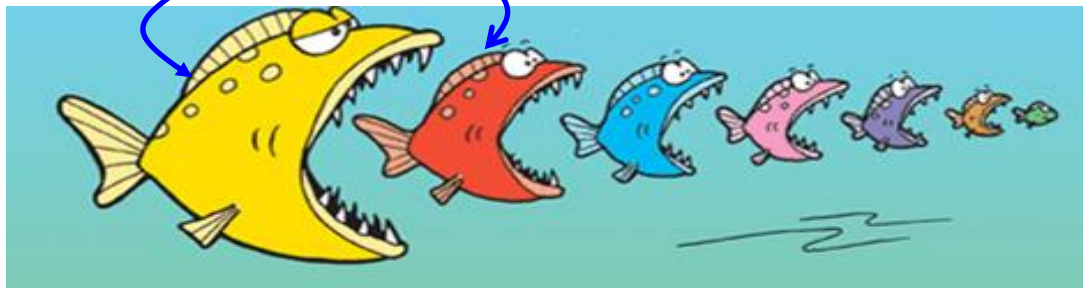
Journal of Archaeological Science (1999) **26**, 667–673
Article No. jasc.1998.0391, available online at <http://www.idealibrary.com>

Isotope Fractionation: Why Aren't We What We Eat?

Dale A. Schoeller

ISOTOPIC DISCRIMINATION IN ECOLOGY

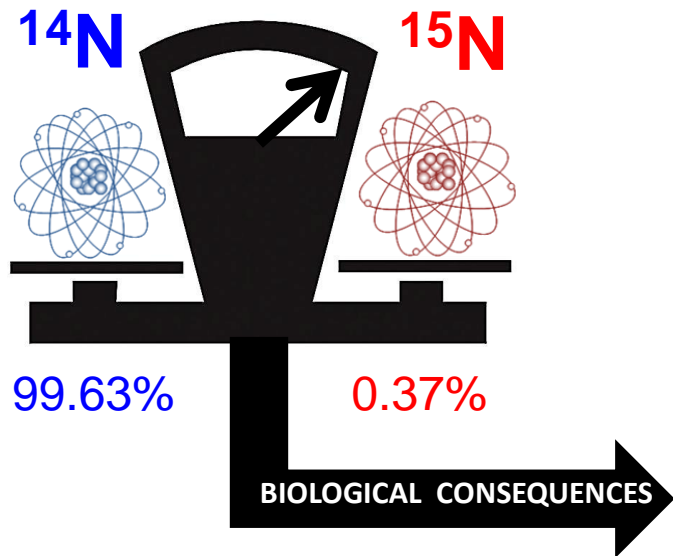
$$\delta^{15}\text{N}_{\text{predator}} - \delta^{15}\text{N}_{\text{prey}} = \Delta^{15}\text{N} = 3.5\text{‰} \pm X$$



$\pm X$ may translate differences in N assimilation across individuals

« Organisms are ^{15}N enriched relative to their diets because not all ingested proteins are retained (some are excreted) and **some metabolic pathways may discriminate ^{15}N vs ^{14}N during the nitrogen assimilation process** »

ISOTOPIC NITROGEN DISCRIMINATION IN RUMINANTS



ENZYMES

HEPATIC AMINO ACID (AA) CATABOLISM :

Affinity for ^{14}N -AA > ^{15}N -AA

(Macko et al., 1986)



- Animal proteins naturally **more enriched in ^{15}N** when AA CATABOLISM increases

(Cantalapiedra-Hijar et al., 2015)

N ASSIMILATION BY RUMEN BACTERIA :

Affinity for ^{14}N - NH_3 > ^{15}N - NH_3

(Wattiaux and Reed, 1995)

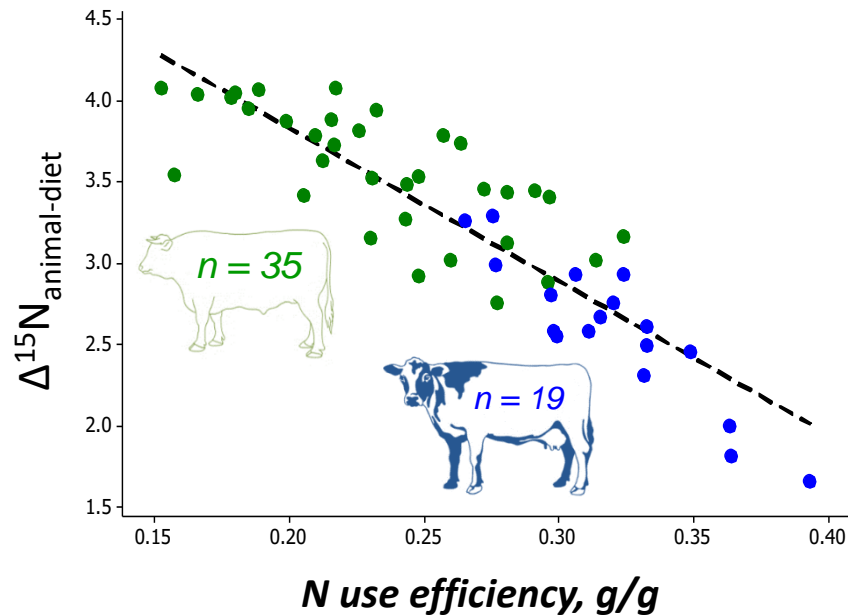


- Animal proteins naturally **more enriched in ^{15}N** when MICROBIAL N ASSIMILATION decreases

(Cantalapiedra-Hijar et al., 2016)

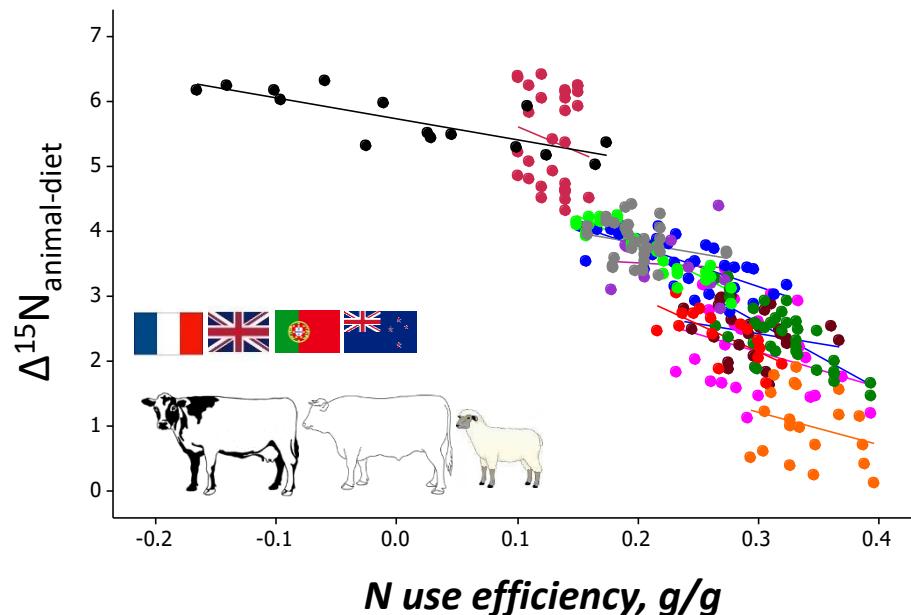
PROTEIN EFFICIENCY AND ISOTOPIC N DISCRIMINATION

PROOF OF CONCEPT



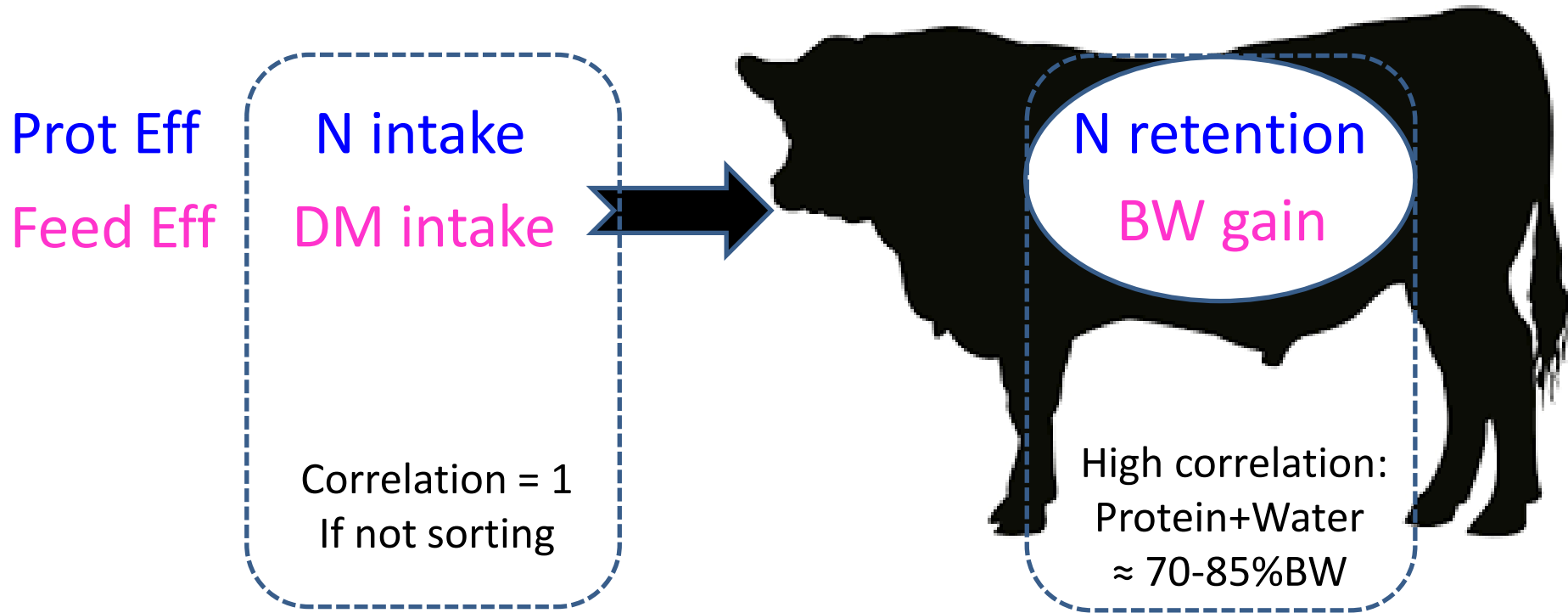
Cantalapiedra-Hijar et al., 2015

CONFIRMATION BY META-ANALYSIS



Cantalapiedra-Hijar et al., 2018

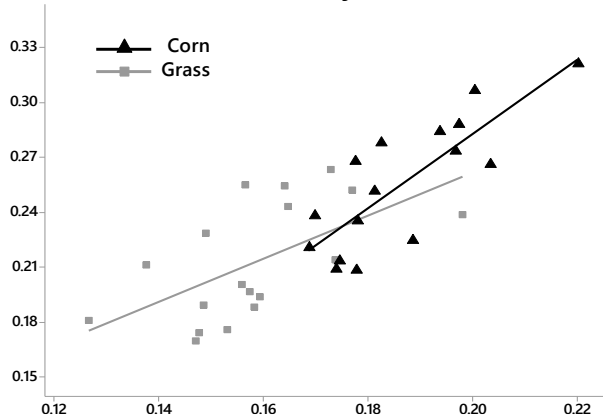
PROTEIN EFFICIENCY AND FEED EFFICIENCY ARE CLOSELY RELATED AT THE INDIVIDUAL LEVEL!!



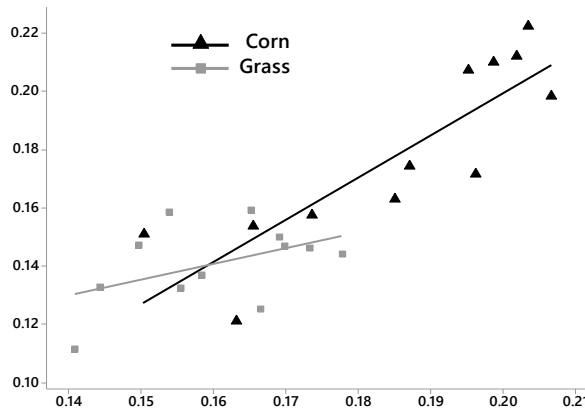
LINK BETWEEN PROTEIN EFFICIENCY AND FEED EFFICIENCY

Protein efficiency, kg/kg

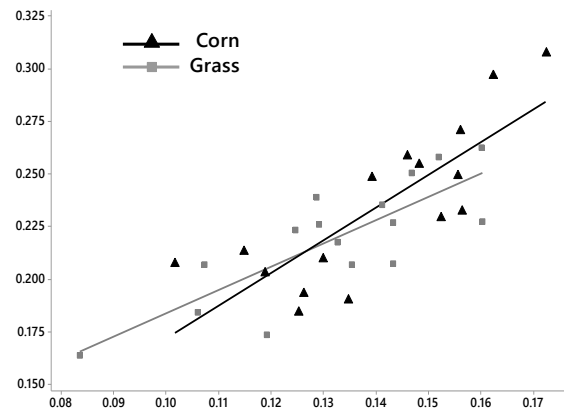
Cantalapiedra-Hijar et al., 2015



Nasrollahi et al., 2020

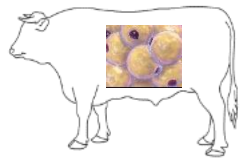


Guarnido et al., unpublished

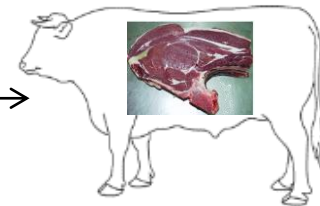


Feed conversion efficiency, kg/kg

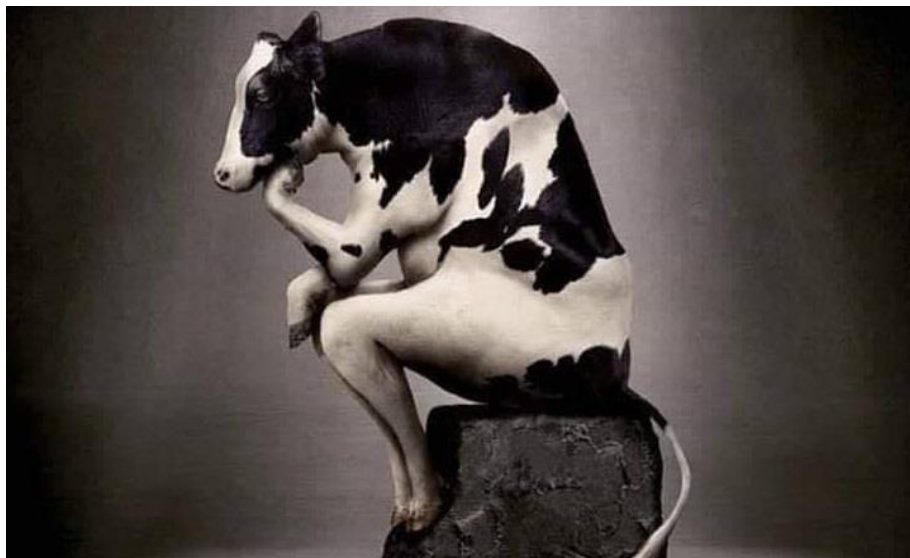
ESTIMATION OF PROTEIN
RETENTION



SUBCUTANEOUS
ADYPOCITES SIZE



6th RIB DISSECTION



Source: <https://edairynews.com/>

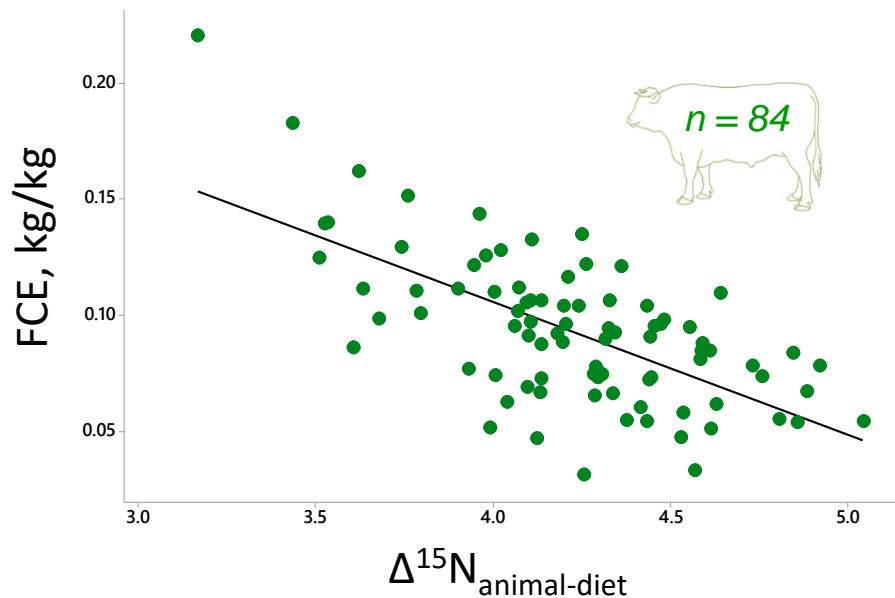
SYLLOGISM:
IF $\Delta^{15}\text{N}$ IS CORRELATED TO PROTEIN
EFFICIENCY

AND PROTEIN EFFICIENCY IS
CORRELATED TO FEED EFFICIENCY

THEN.....

FEED EFFICIENCY AND $\Delta^{15}\text{N}$

PROOF OF CONCEPT



Wheardon et al., 2014

CONFIRMED IN SOME MORE STUDIES

- Cantalapiedra-Hijar et al., 2015
- Meale et al., 2017
- Meale et al., 2018
- Nasrollahi et al., 2020
- Cantalapiedra-Hijar et al., 2020

INCONSISTENT WITH RFI

- Non-Significant:
 - Wheardon et al., 2014;
 - Meale et al., 2017
- Significant:
 - Cantalapiedra-Hijar et al., 2020



OBJECTIVES

To confirm by meta-analysis the potential of $\Delta^{15}\text{N}_{\text{animal-diet}}$ to reflect animal-to-animal variation of feed conversion efficiency in different EU beef production systems

*To test if this biomarker could equally work for residual feed intake (not originally included in the EAAP abstract)

MATERIAL AND METHODS (I)

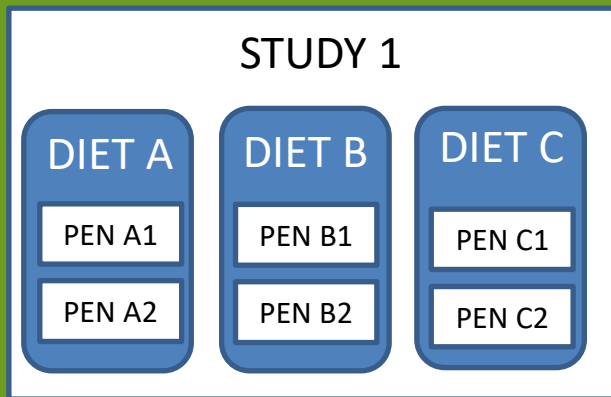
- 9* studies from 3 countries (FR, UK, CH) with FE test lasting from 60 to 200d
Wheadon et al., 2014; Cantalapiedra-Hijar et al., 2015, 2020; Meale et al., 2017,2018; Nasrollahi et al., 2020
+ 3 unpublished studies
- 570* individual data from growing-fattening bulls (6), steers (2) or heifers (1)
- Pure Charolais (25%) or crossed breeds (Lim x Sim, Sim x Hol, Ang x Lim..)
- 27* different fattening diets (9 to 84 individuals per diet)
 - 65% data from diets based on grass silage (>50%DM)
- Feed pooled for the whole FE test and blood/muscle sampling at the end
- Plasma, muscle and feed samples analyzed for $\delta^{15}\text{N}$ by EA-irms

** One more study has been added to this analysis in relation to the EAAP abstract*

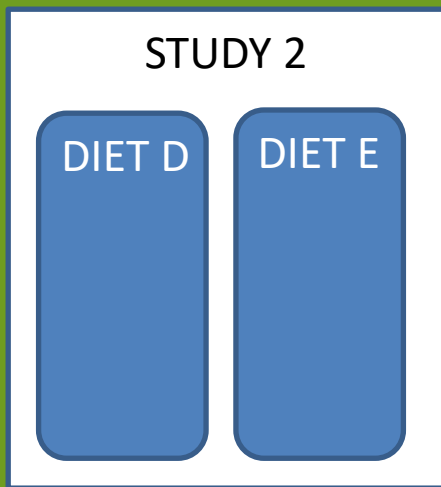
MATERIAL AND METHODS (II)

What we consider to be « at the individual level » in this study ?

STUDY 1



STUDY 2



Animals reared in similar conditions:

- Same study (location)
- Same diet at the same time
- But NOT the same pen (info not available yet)

At the individual level = **Within-DIET regressions** => No need ^{15}N from diet!

MATERIAL AND METHODS (III)

2 statistical approaches to explore relationships at the individual level:

- **I) MIXED-EFFECT MODEL**

$$FCE = (A + \alpha) + (B + \beta) \times \Delta^{15}\text{N} + \varepsilon$$

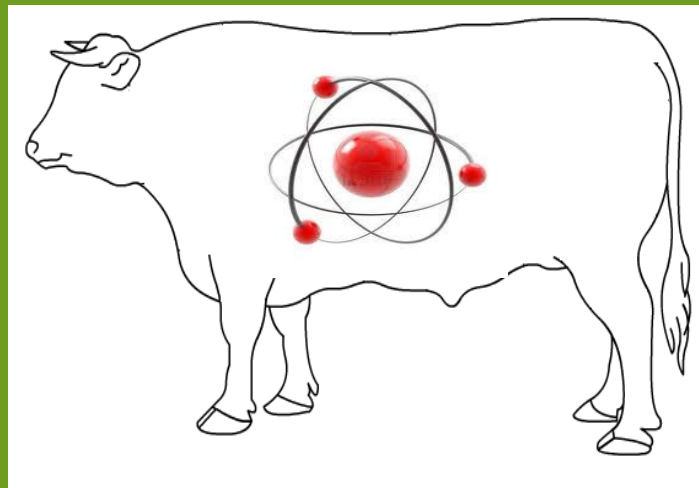
α and β : random effects (study and diet within study)

- **II) SIMPLE REGRESSION** when the effect of study and diets were first removed from both FE and $\Delta^{15}\text{N}$

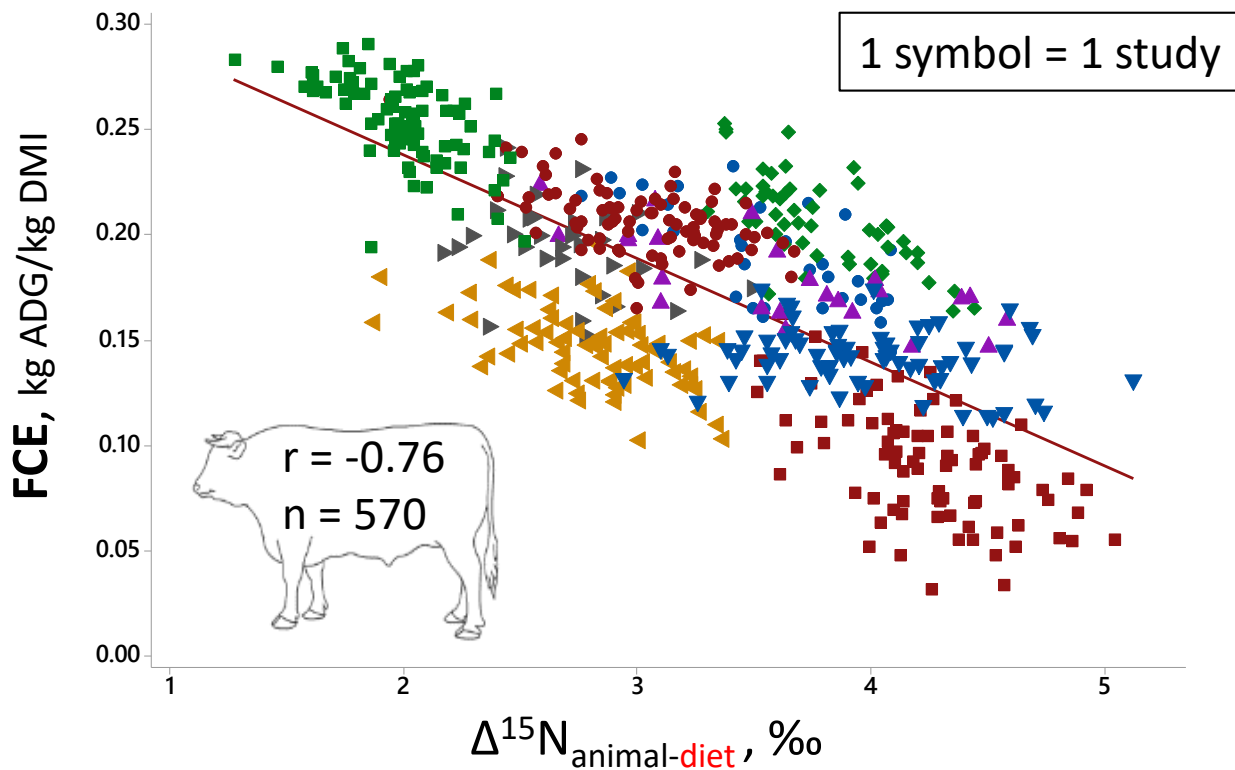
a) Residuals $FCE = A + B \times \text{Residuals } \Delta^{15}\text{N}$

b) $RFI = A + B \times \text{Residuals } \Delta^{15}\text{N}$

RESULTS



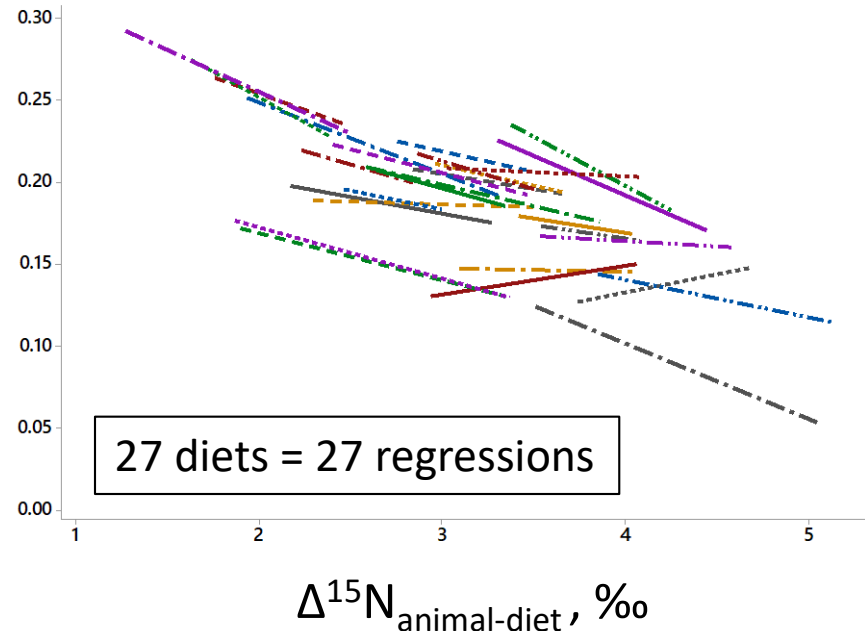
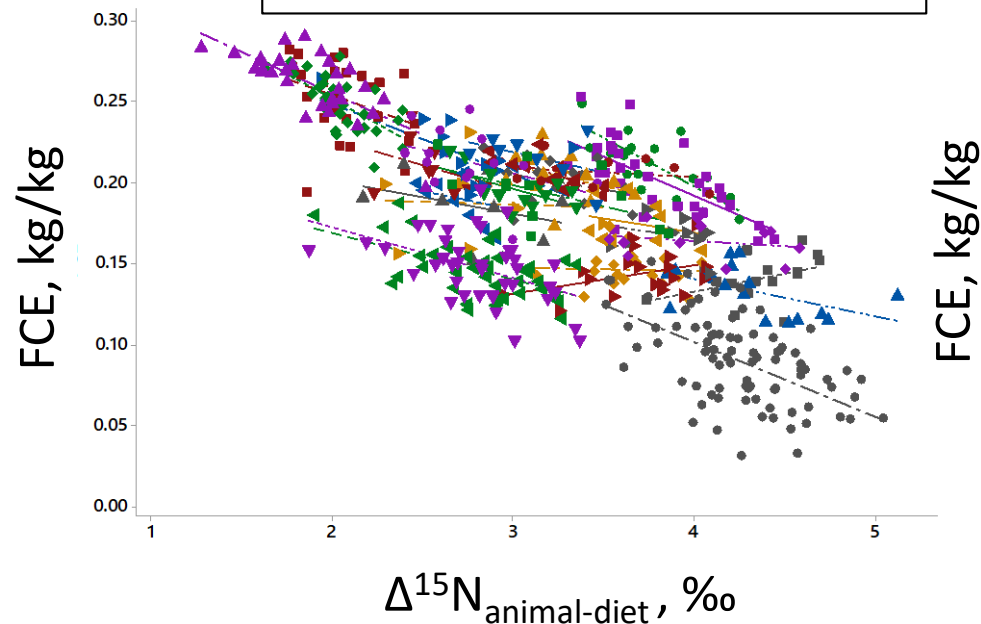
OVERALL RELATIONSHIP BETWEEN FCE & $\Delta^{15}\text{N}$



WITHIN-DIET RELATIONSHIP BETWEEN FCE & $\Delta^{15}\text{N}$

(MIXED-EFFECT MODEL)

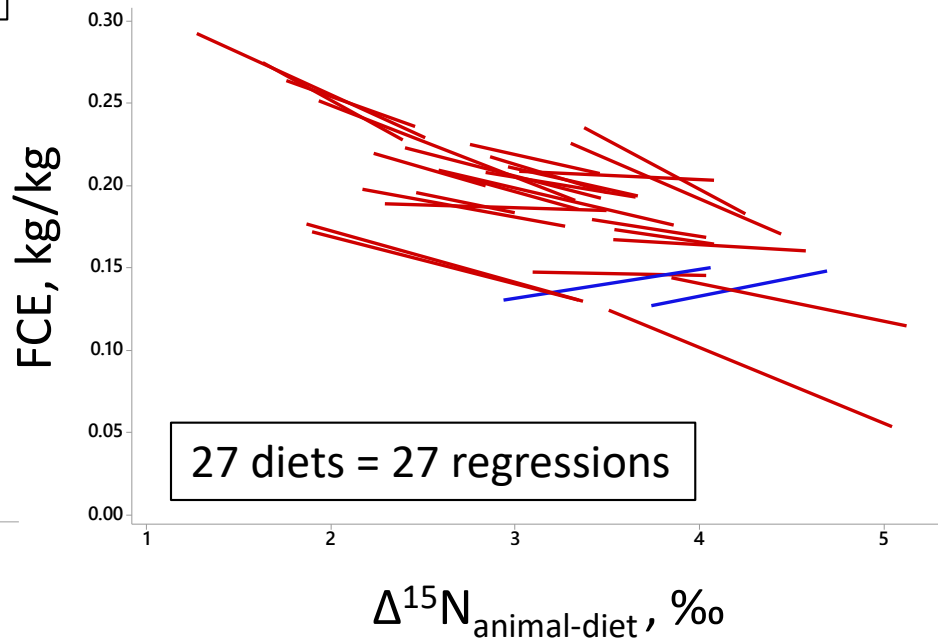
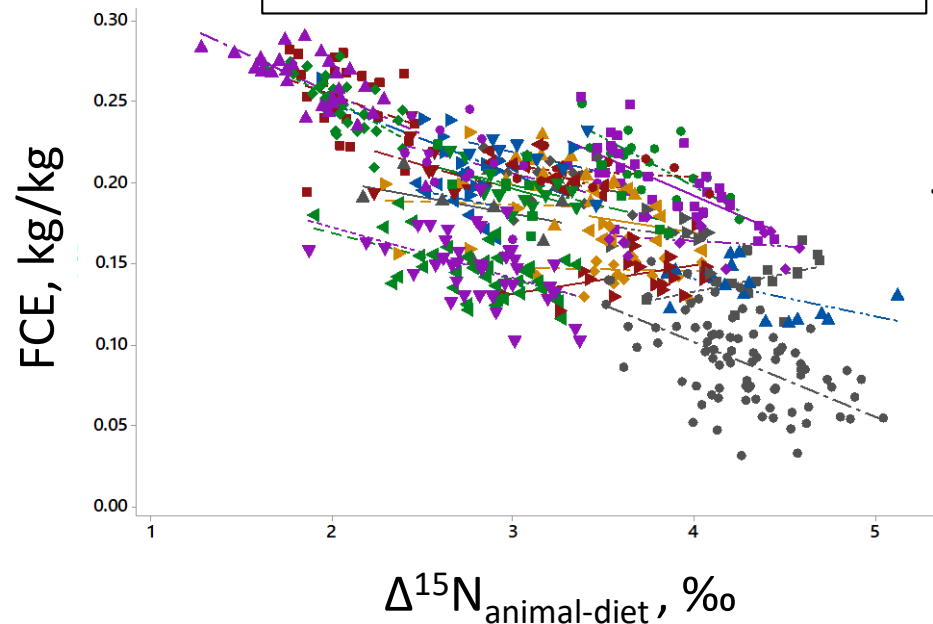
1 symbol = 1 diet within 1 study



WITHIN-DIET RELATIONSHIP BETWEEN FCE & $\Delta^{15}\text{N}$

(MIXED-EFFECT MODEL)

1 symbol = 1 diet within 1 study



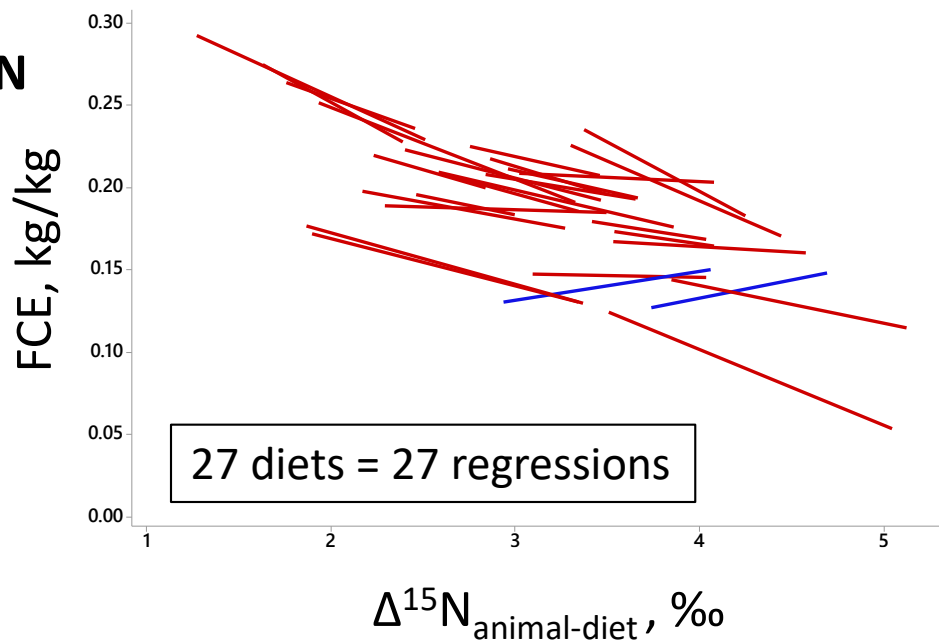
WITHIN-DIET RELATIONSHIP BETWEEN FCE & $\Delta^{15}\text{N}$ (MIXED-EFFECT MODEL)

Average within-diet regression

$$\text{FCE} = 0.28^*(\pm 0.02) - 0.032^{**}(\pm 0.004) \times \Delta^{15}\text{N}$$

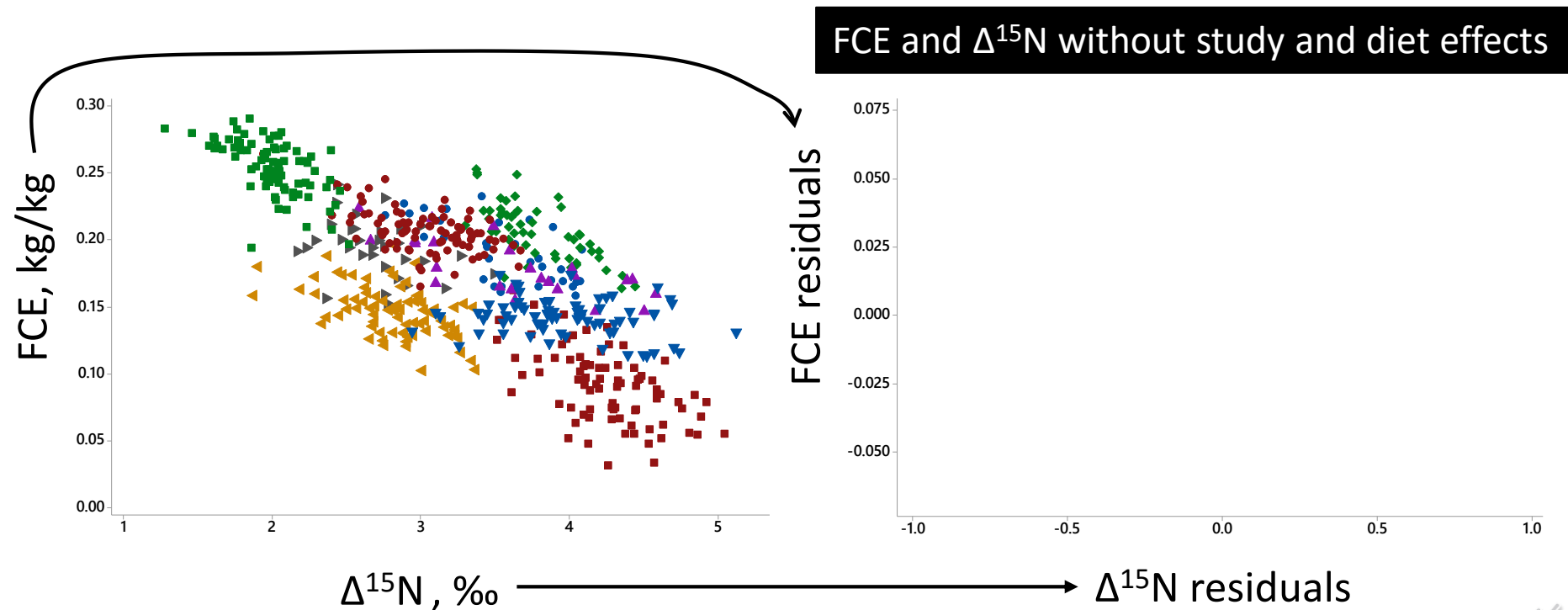
$$\text{RSE} = 0.017 \text{ kg/kg}$$

😊 The error is still compatible with the identification of extreme FCE young bulls!



WITHIN-DIET RELATIONSHIP BETWEEN FCE & $\Delta^{15}\text{N}$

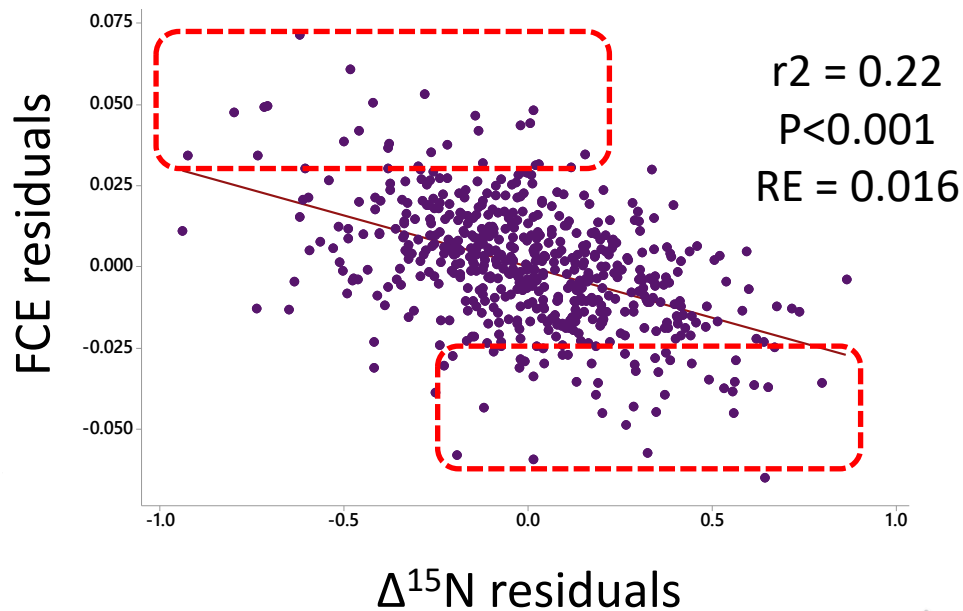
(SIMPLE REGRESSION FROM RESIDUALS)



WITHIN-DIET RELATIONSHIP BETWEEN FCE & $\Delta^{15}\text{N}$

(SIMPLE REGRESSION FROM RESIDUALS)

FCE and $\Delta^{15}\text{N}$ without study and diet effects

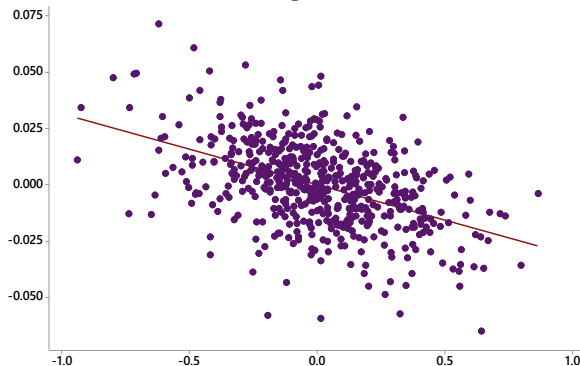


WITHIN-DIET RELATIONSHIP BETWEEN PERFORMANCES AND $\Delta^{15}\text{N}$

(SIMPLE REGRESSION FROM RESIDUALS)

FCE

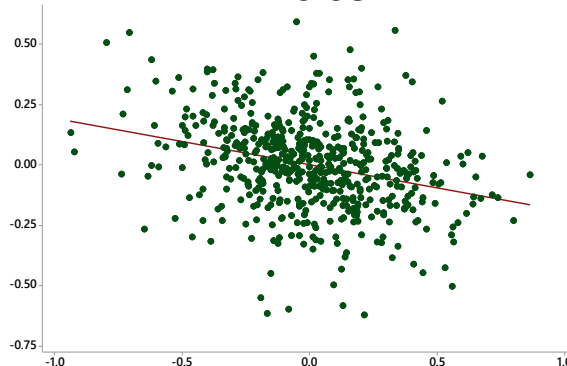
$r^2 = 0.22^{***}$



$\Delta^{15}\text{N}$ residuals

ADG

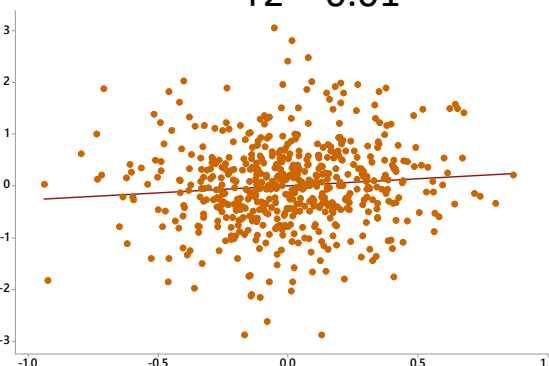
$r^2 = 0.08^{***}$



$\Delta^{15}\text{N}$ residuals

DMI

$r^2 = 0.01^*$



$\Delta^{15}\text{N}$ residuals

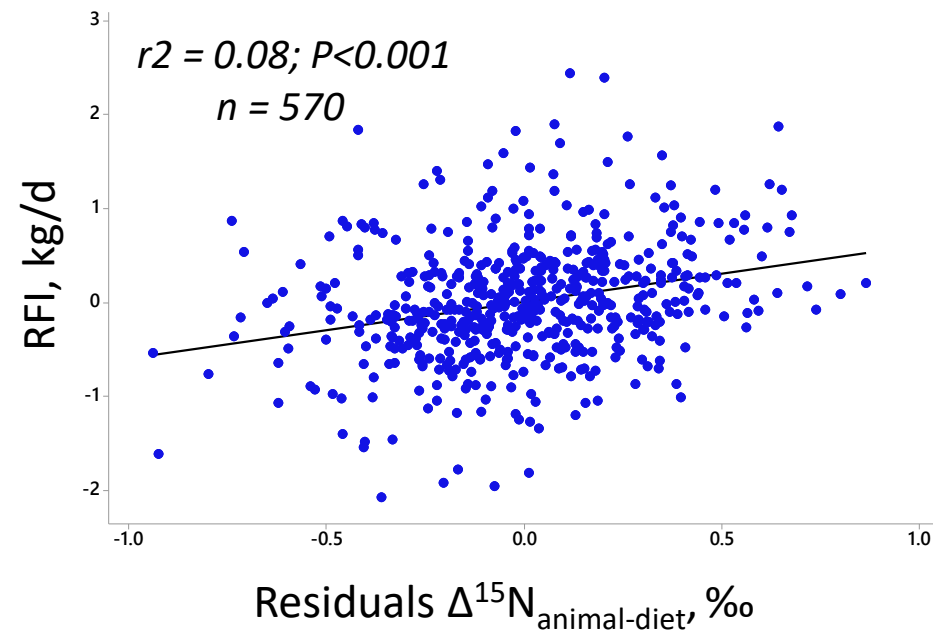
FCE \sim ^{15}N ($r^2 = 0.22$)
FCE \sim ADG ($r^2 = 0.44$)

If we combine both predictors: FCE \sim ADG + ^{15}N ($r^2 = 0.52$)

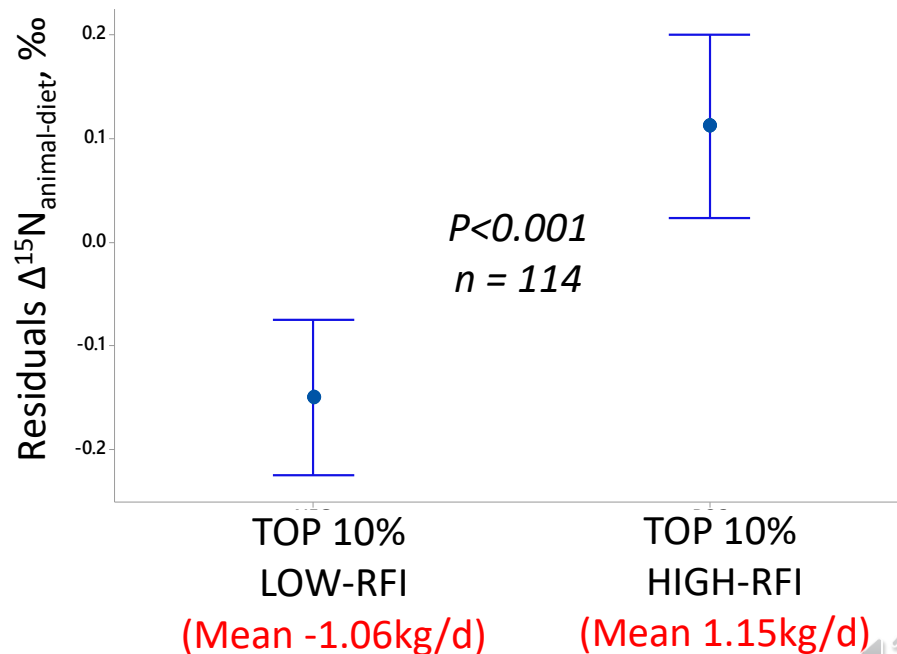
.....but need to be validated

POTENTIAL OF $\Delta^{15}\text{N}$ TO REFLECT RFI VARIATION

WHOLE POPULATION



EXTREME RFI ANIMALS



CONCLUSIONS

- We confirmed the potential of $\Delta^{15}\text{N}$ to reflect between-animal variation in feed conversion efficiency and residual feed intake
- Most efficient beef cattle (10% highest FCE and 10% lowest RFI) had lower ^{15}N abundance in their proteins than their less efficient counterparts
- More research is needed to validate this biomarker in practical conditions and to assess genetic correlations with feed efficiency

ACKNOWLEDGEMENTS



I Ortigues-Marty



C Martin



I Morel



R Dewhurst



Céline Chantelauze
(irms analysis)

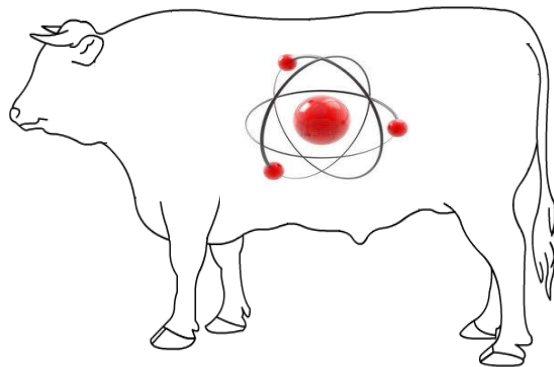
Public funds



Private funds



THANK YOU FOR YOUR ATTENTION



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