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# IMPROVED CATTLE GROWTH BY METHIONINE- BALANCED DIETS DOES NOT RESULT FROM LOWER PROTEIN DEGRADATION

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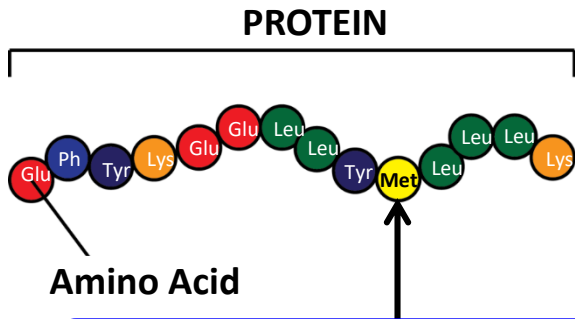




# INTRODUCTION

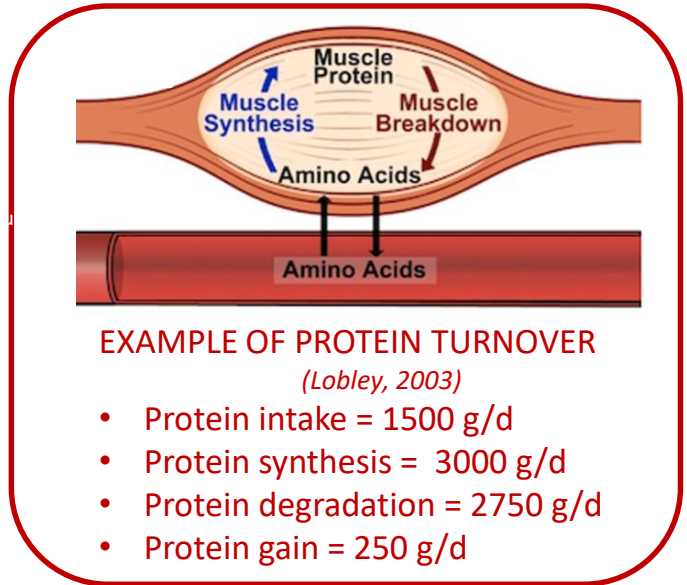
- Improving N use efficiency (NUE) is currently an important issue in animal production
- From a metabolic point of view both the amount and nature of protein may impact NUE
- The dietary AA profile should be formulated according to animal requirements
  - Methionine is believed to be the first limiting AA in growing cattle fed forage-diets (*Titgemeyer and Merche, 1990*)
  - Diets well balanced for Met improve performances of growing beef cattle (*Veira et al., 1991; Bahloul et al., 2018*)
  - However, controversy exists about the metabolic pathways involved in this improvement

# PROTEIN TURNOVER

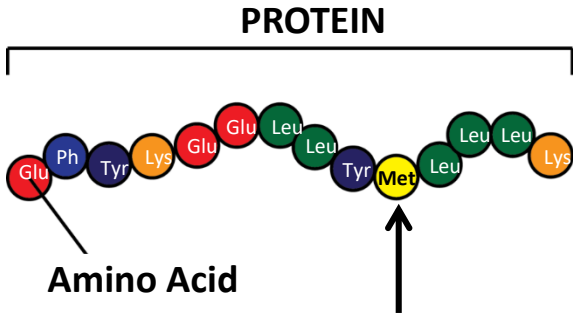


**IF THIS AMINO ACID LACKS:**

- 1) PROTEIN SYNTHESIS MAY BE REDUCED**
- 2) PROTEIN DEGRADATION MAY INCREASE**

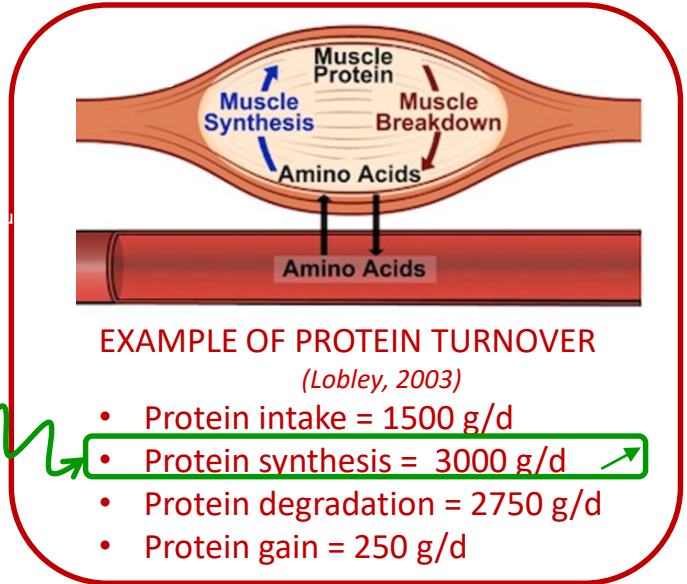


# PROTEIN TURNOVER: H1

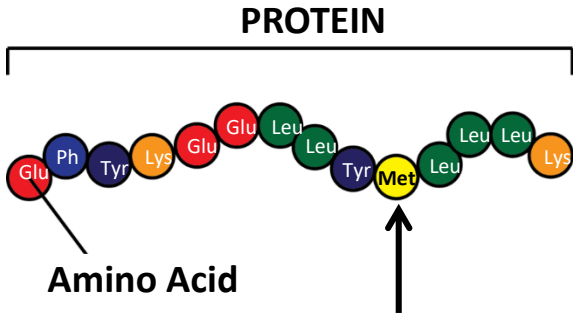


**IF THIS AMINO ACID IS SUPPLEMENTED  
THE PROTEIN DEPOSITION MAY  
INCREASE THROUGH:**

*Salter et al., 1990; Wessels et al., 1997;  
Saggau et al., 2000; Ren et al., 2007  
Robinson et al., 2016*

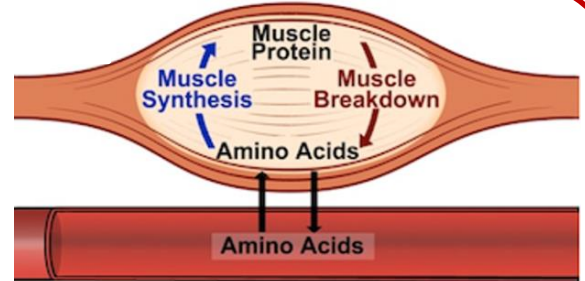


# PROTEIN TURNOVER: H2



**IF THIS AMINO ACID IS SUPPLEMENTED  
THE PROTEIN DEPOSITION MAY  
INCREASE THROUGH:**

*Nieto et al., 1994; Tesseraud et al., 1996  
De la Higuera et al., 1998; Schadereit et al., 1999  
Löhcke et al., 2001*



## EXAMPLE OF PROTEIN TURNOVER

*(Lobley, 2003)*

- Protein intake = 1500 g/d
- Protein synthesis = 3000 g/d
- Protein degradation = 2750 g/d
- Protein gain = 250 g/d



**Mechanisms responsible for the effect of Met (synthesis vs degradation) are not elucidated.**

**Available studies:**

- ✓ **monogastrics only (pigs, broilers, rats, fish)**
- ✓ **measurement of protein synthesis using a reference method (infusion or flooding dose of a labelled amino acid)**
- ✓ **but no measurement of protein degradation (→ calculated).**

## **OBJECTIVE**

**To analyse the whole-body protein turnover rate of fattening young bulls fed diets balanced or unbalanced for methionine, at two dietary levels of metabolizable protein, using a new methodology to quantitatively assess protein degradation rate in vivo.**

# MATERIAL AND METHODS

- 36 Charolais young bulls (320 kg BW and 266 d old on average)
- 4 experimental diets, all based on grass silage (60%) and concentrate (40%)

## 2 x 2 Factorial design

[Normal vs High MP] x [Without vs with Smartamine®]  
100 vs 120% requirements x 1.9 vs 2.4 %Met (Lys/Met ~ 4 vs 3)

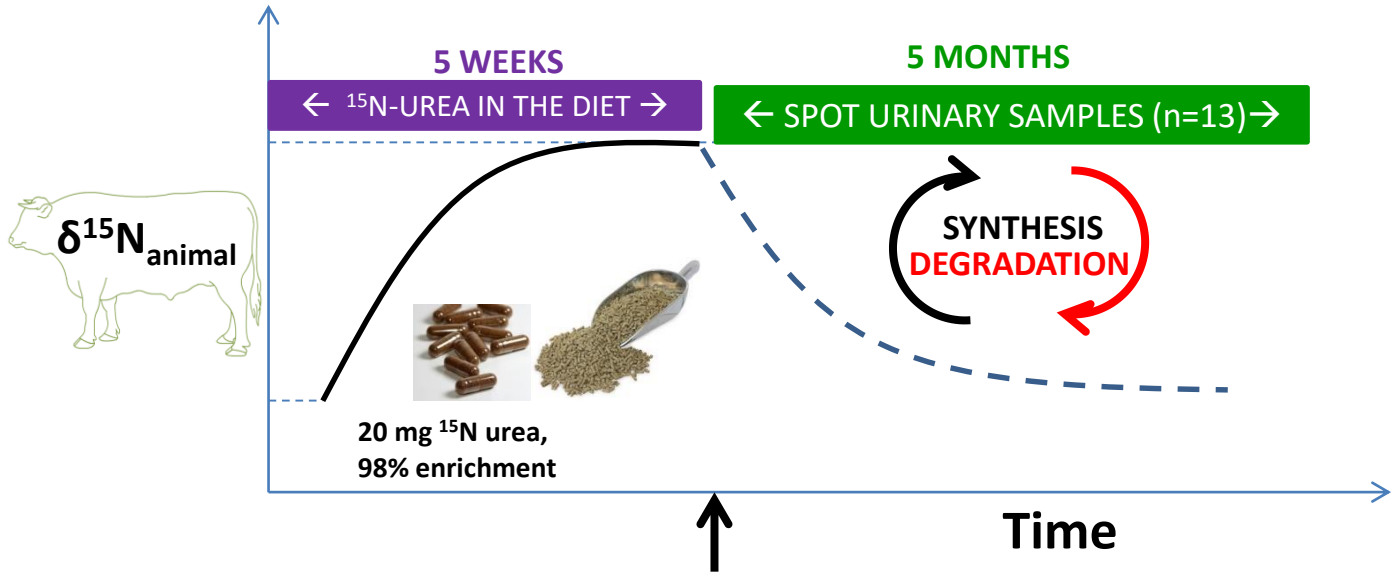
The 2 experimental factors (MP and Met) significantly increased ADG (+16 and 9%, respectively) (*Bahloul et al., 2018*) and thus were supposed to impact the protein metabolism

## Measurement of isotopic ( $^{15}\text{N}$ ) turnover

- ✓ *After tissue enrichment in  $^{15}\text{N}$ , the rate of release of  $^{15}\text{N}$  from the whole body reflects WB protein degradation*
- ✓ *≠ reference methods which target protein synthesis using tracers*

# ISOTOPIC $^{15}\text{N}$ TURNOVER RATE

Measurement in urine following an isotopic diet switch



The rate at which WB proteins release  $^{15}\text{N}$  after accumulation reflects WB protein degradation rate.

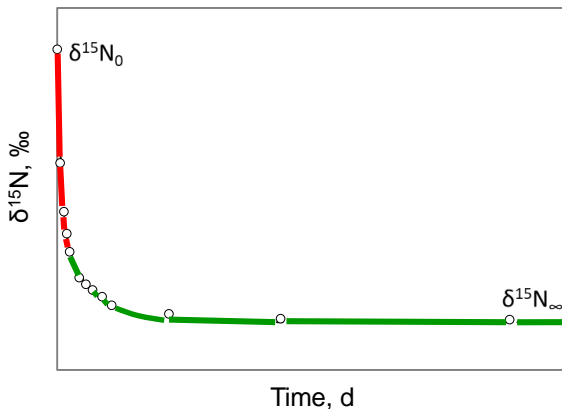
Protein synthesis rate evaluated by difference from ADG and protein degradation rate.



# MODELING of ISOTOPIC TURNOVER RATE

$$\delta^{15}\text{N}(t) = \delta^{15}\text{N}_{\infty} + (\delta^{15}\text{N}_0 - \delta^{15}\text{N}_{\infty}) \times [p \times \exp^{-k_1 \times t} + (1-p) \times \exp^{-k_2 \times t}]$$

Slope  $k_1$  = degradation rate of pool 1 (fast)  
Slope  $k_2$  = degradation rate of pool 2 (slow)



- All individual data are used to adjust a non linear mixed-effect model (nlme in R)  
Mono or bi-exponential
- Fixed effects : MP level, Methionine content and their interaction
- Random effect: Animal



# RESULTS

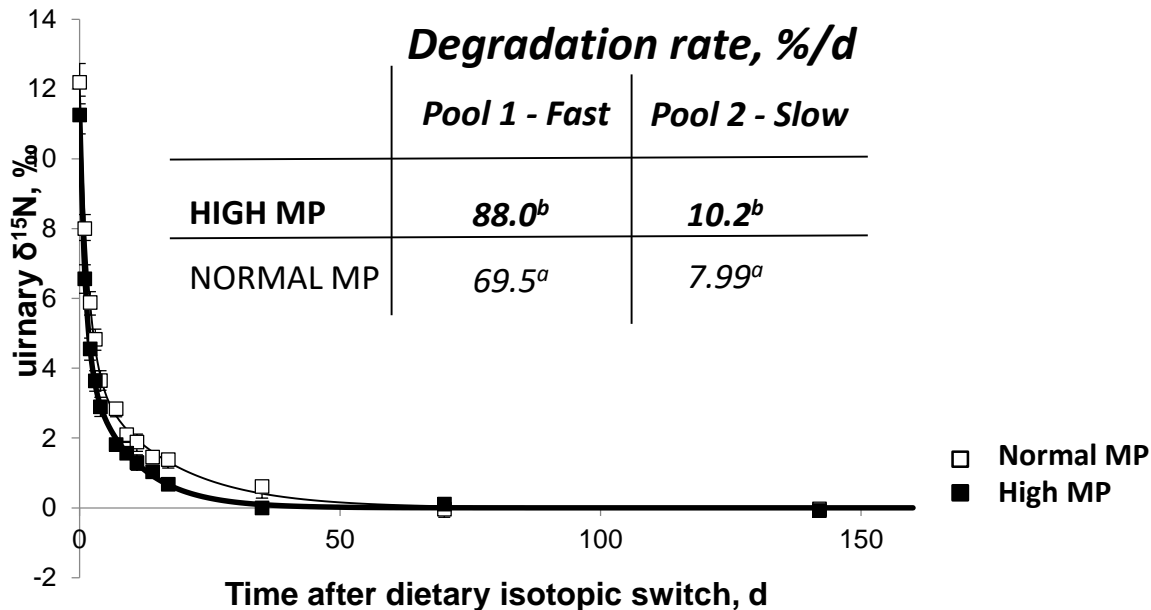
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# Effect of DIETARY PROTEIN LEVEL



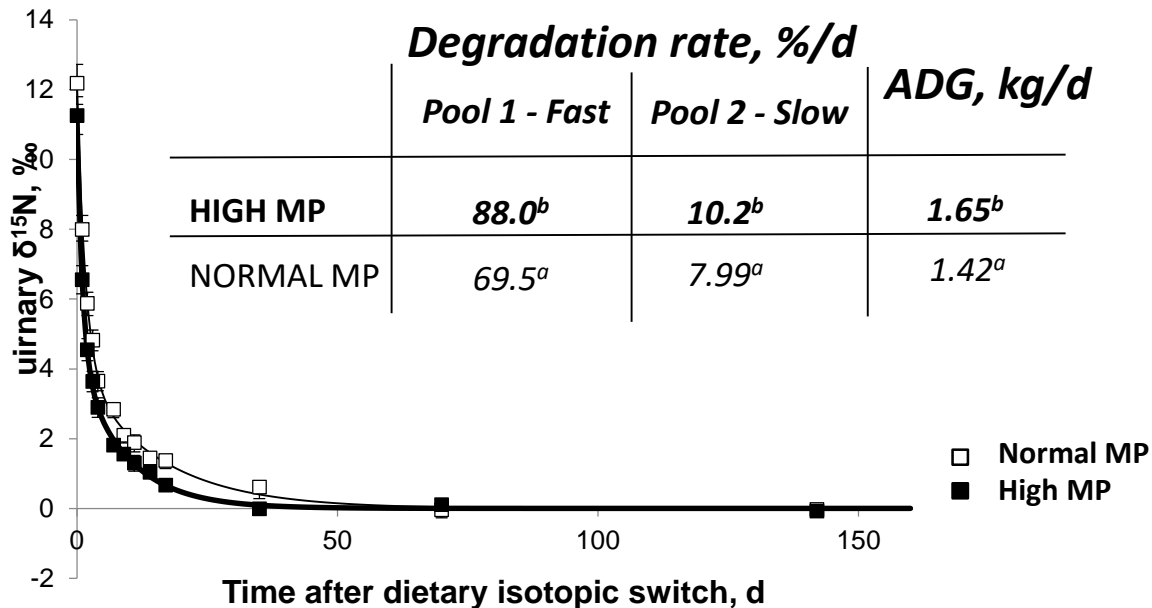
# Effect of DIETARY PROTEIN LEVEL

## High MP increases the whole body protein degradation rate



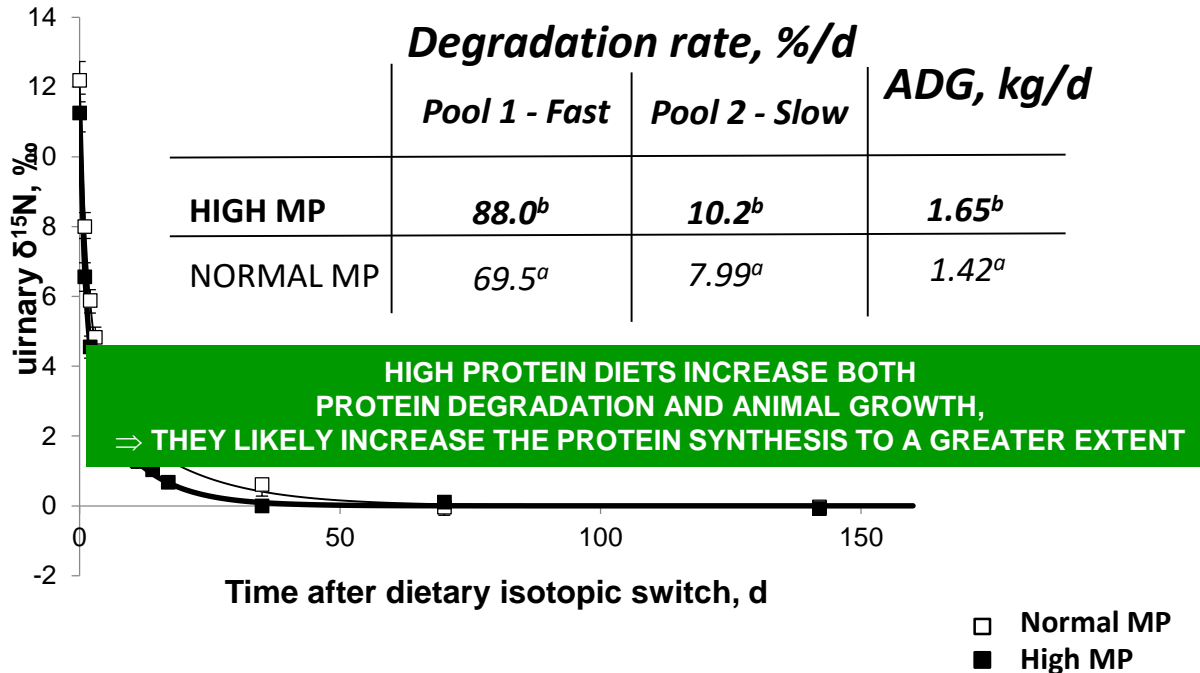
# Effect of DIETARY PROTEIN LEVEL

High MP increases the whole body protein degradation rate



# Effect of DIETARY PROTEIN LEVEL

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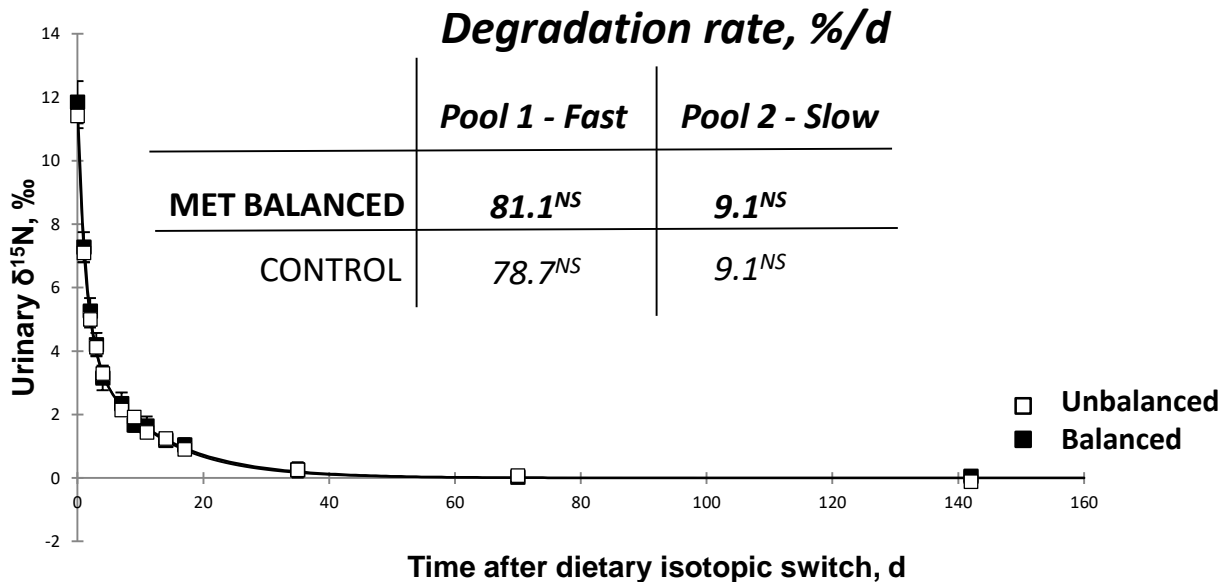


# Effect of DIETARY METHIONINE LEVEL



# Effect of DIETARY METHIONINE LEVEL

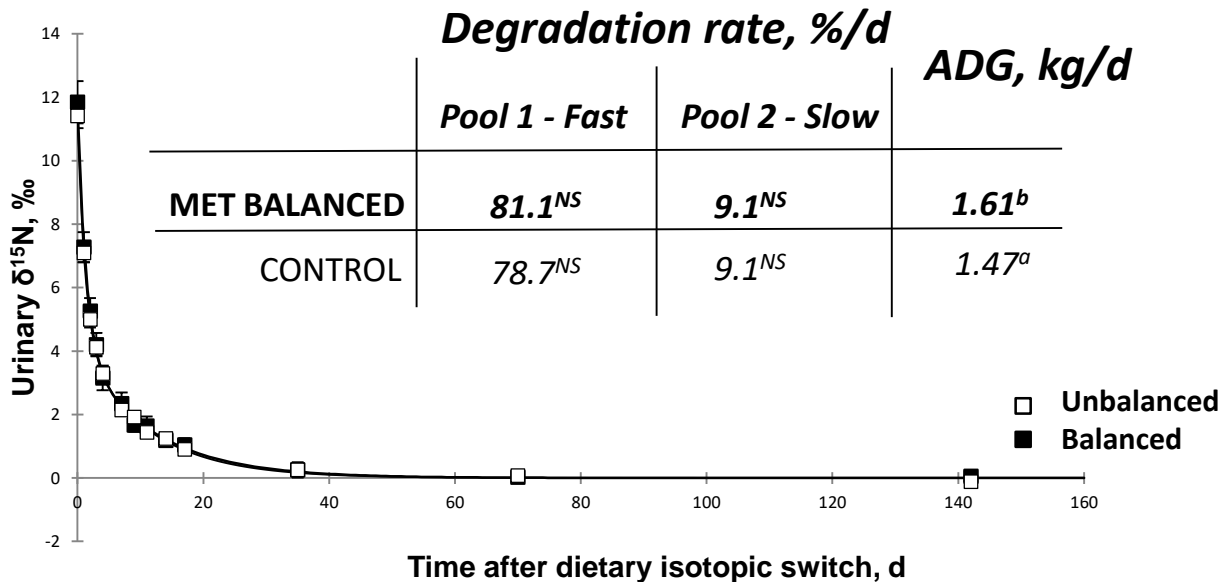
## No impact on whole body protein degradation





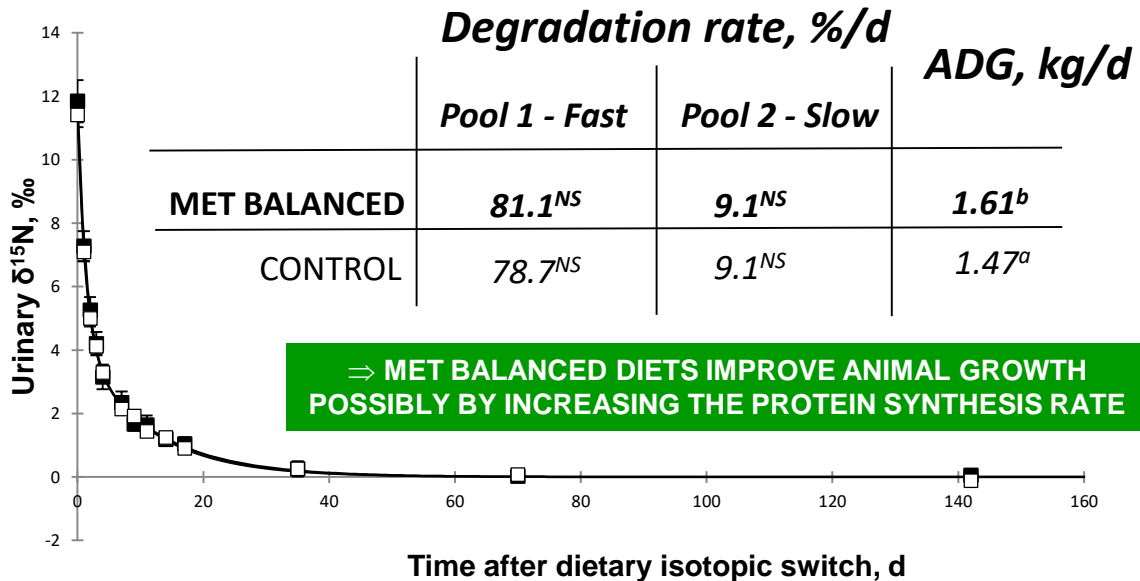
# Effect of DIETARY METHIONINE LEVEL

## No impact on whole body protein degradation



# Effect of DIETARY METHIONINE LEVEL

## No impact on whole body protein degradation



□ Unbalanced  
 ■ Balanced



# CONCLUSIONS

The improvement of animal growth with methionine balanced diets is not due to a decrease in whole-body protein degradation rate but more likely to an increase in protein synthesis.

As expected, increasing the protein content of diets increased the whole-body protein degradation rate, and may also have increased the protein synthesis rate.