

A new methodology to estimate protein feed value using the milk protein biological response

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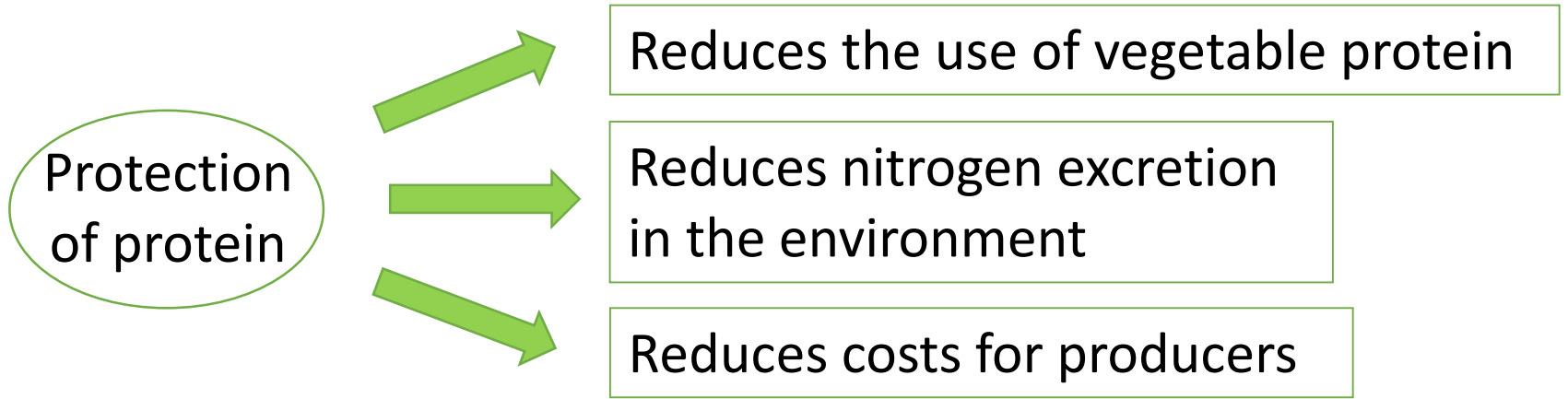
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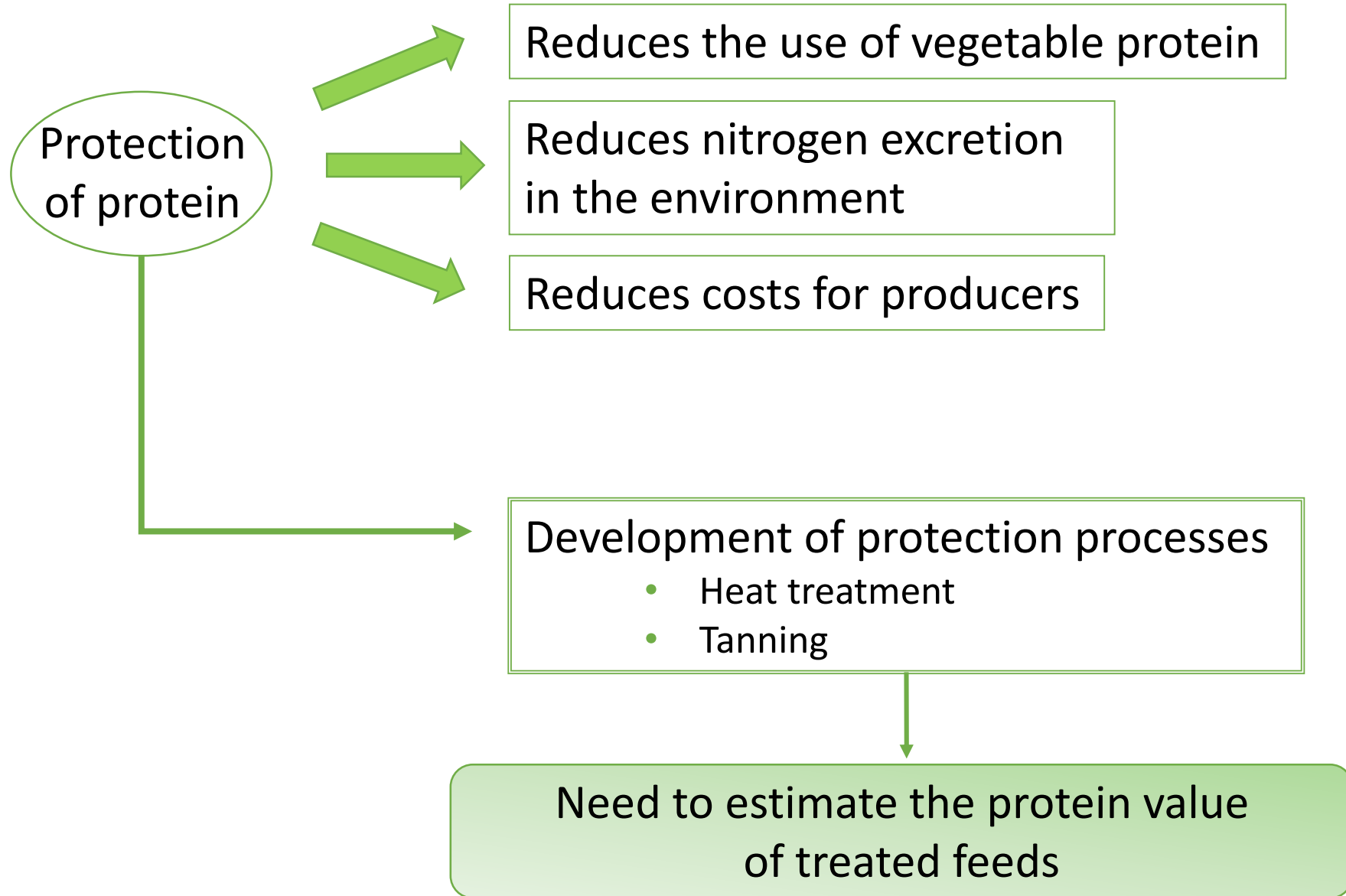
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Introduction



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- Methods used to estimate protein value of protected feeds
 - *In vivo*: measurement of nitrogen duodenal flow
 - Fistulated cows
 - Low accuracy, expensive and time-consuming
 - *In vitro*: test of degradability with ruminal extracted or commercial enzymes
 - Large number of samples tested
 - Bias links to selection of enzymes
 - *In sacco*: test of degradability of feeds by incubation in rumen
 - Most commonly used
 - Particulate losses and microbial colonization in bags

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 - *In sacco*: test of degradability of feeds by incubation in rumen
 - Most commonly used
 - Particulate losses and microbial colonization in bags
- Development of new techniques to protect protein from ruminal fermentations (essential oils, vegetable tannins)
 - Have systemic effects in the rumen
 - Can not be estimated with classical methods

Introduction

- Need for a new and more systemic method to estimate feeds protein value
- Hypothesis: biological response can be used as new method
 - Overall response of animal to protein intake
 - Comparison of different techniques of protein protection

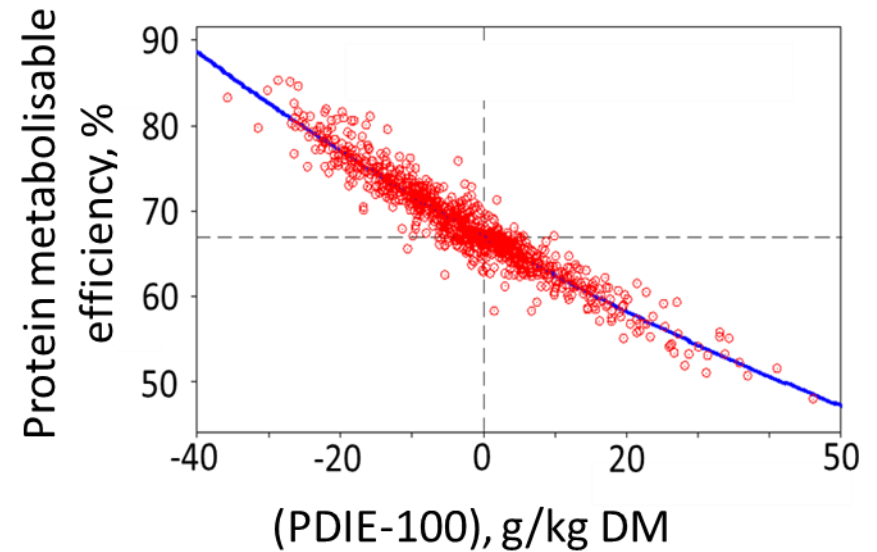
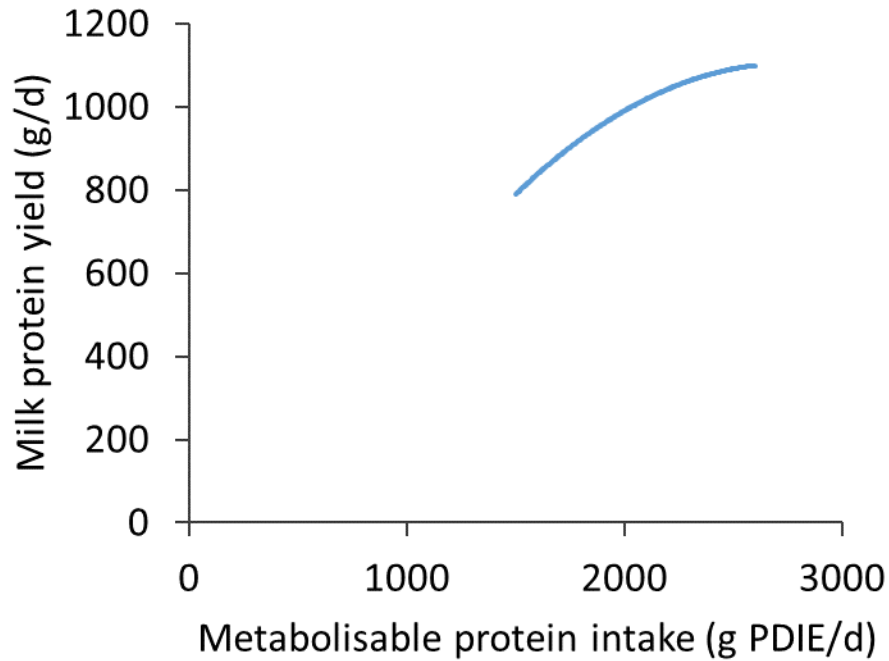
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Can milk protein yield response can be used as a new method to estimate feeds protein value?

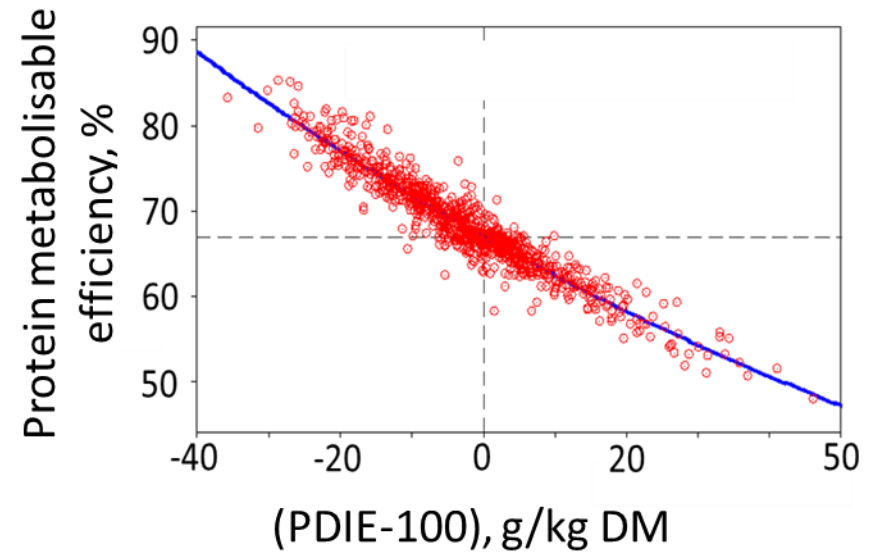
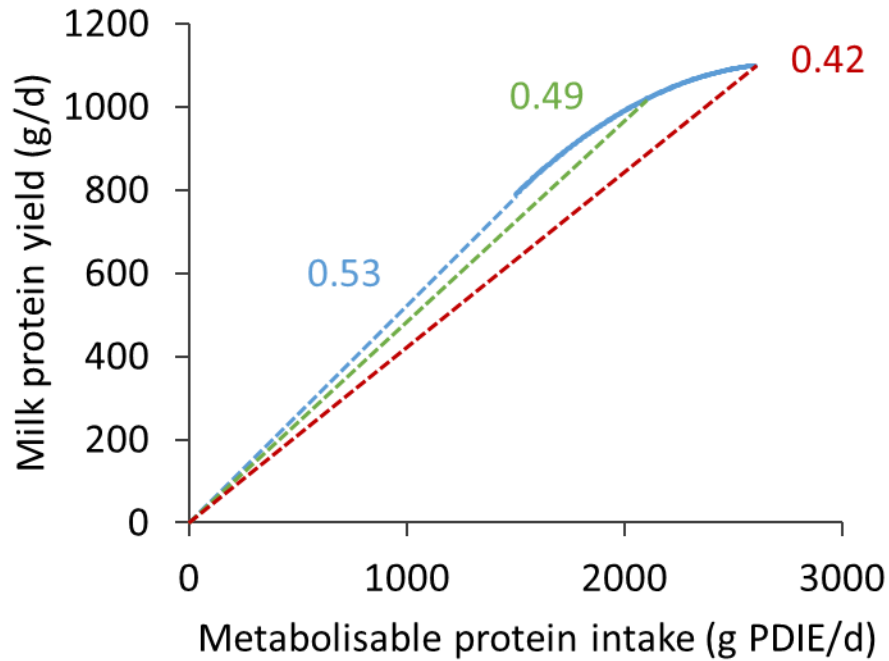
- Protein value of an unknown feed can be determined from its milk protein yield response
- Use of two known dietary controls (positive and negative)

Principle of the method



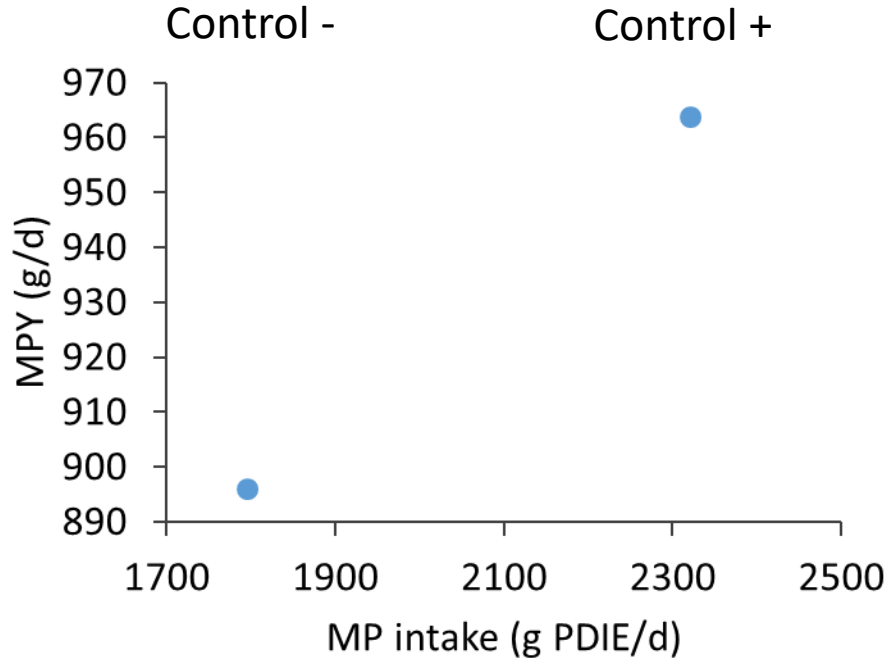
(Sauvant et al. 2015)

Principle of the method



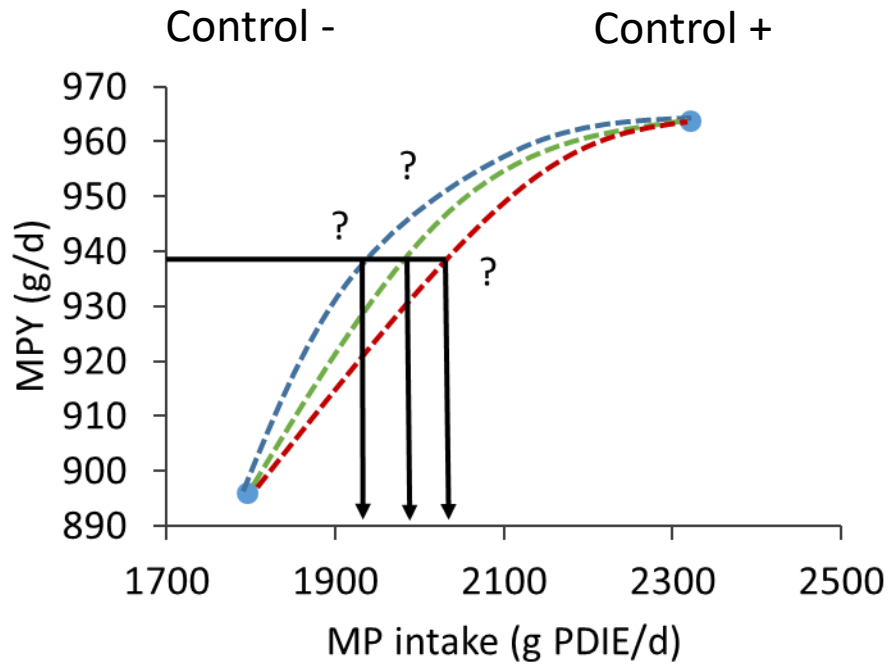
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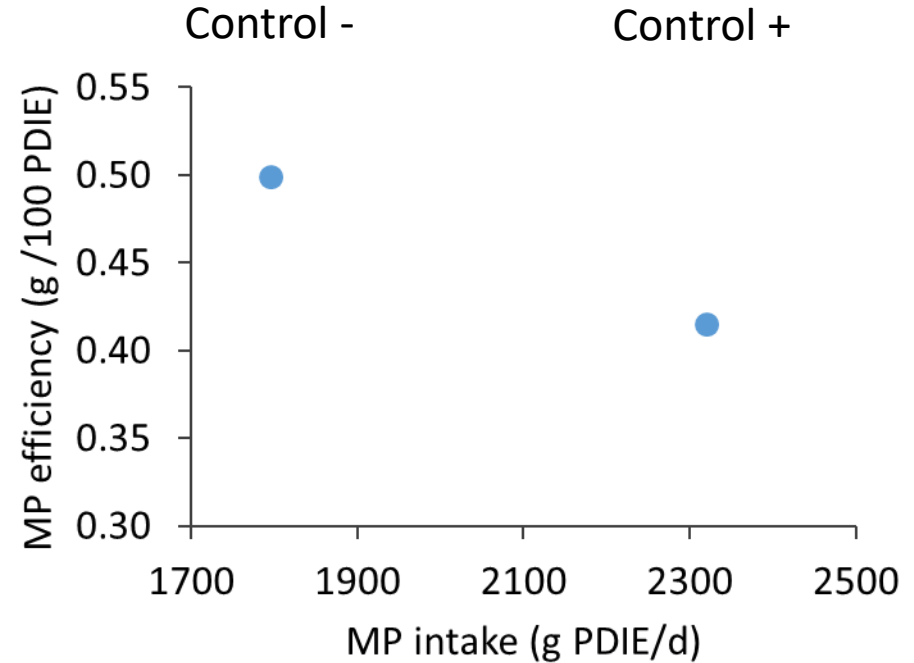
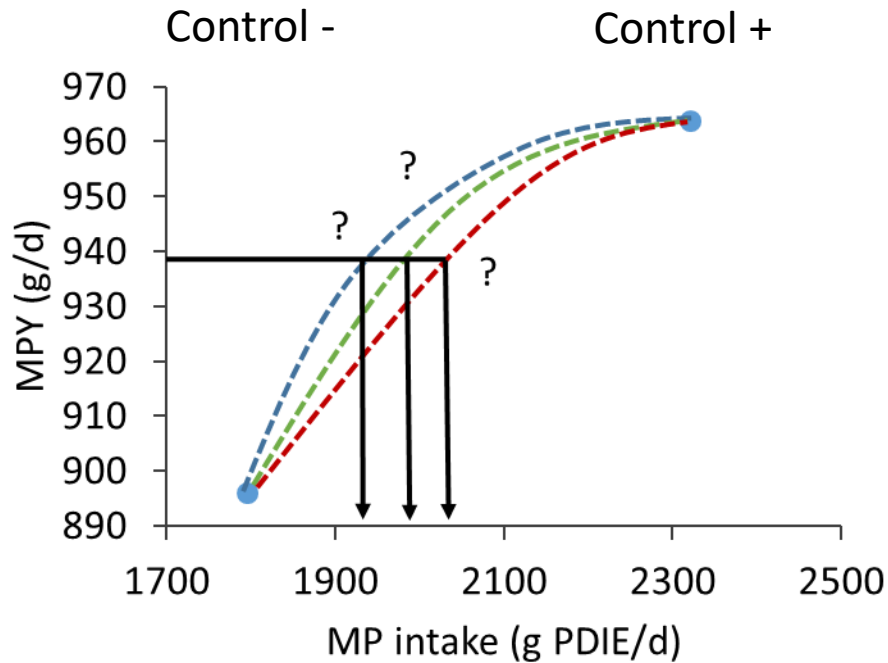
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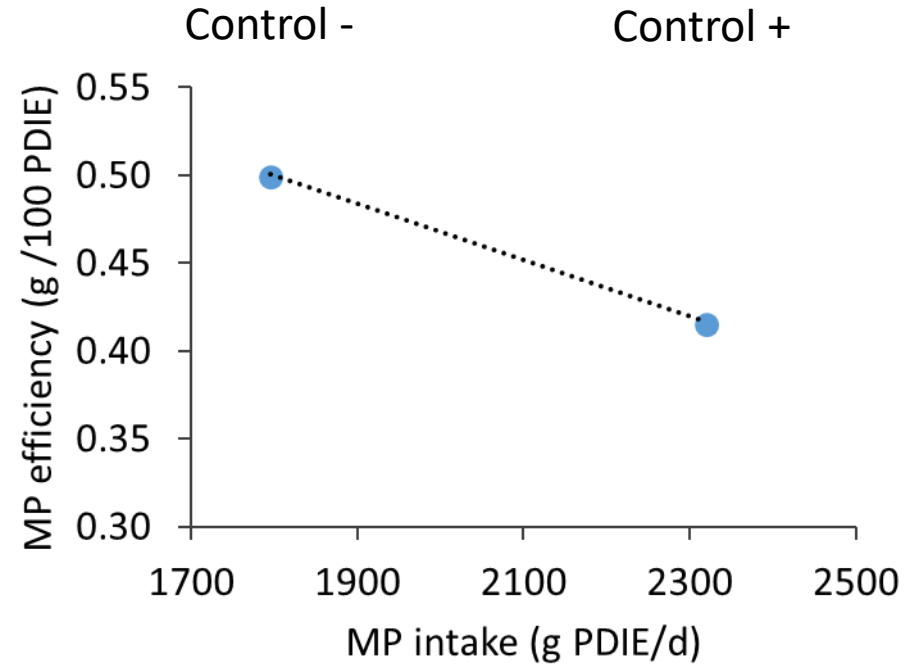
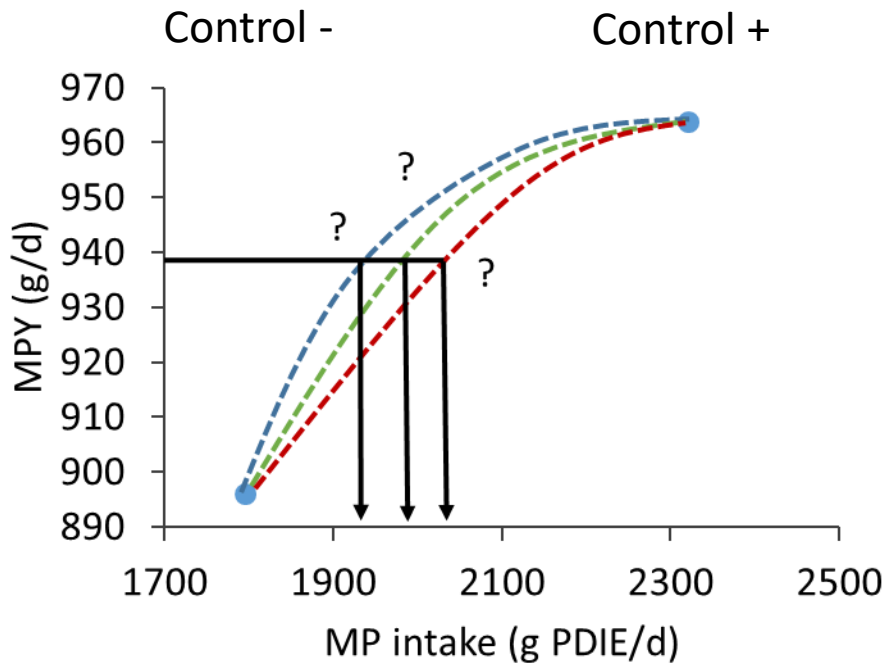
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- $MP\ eff = MPY / MP\ intake$

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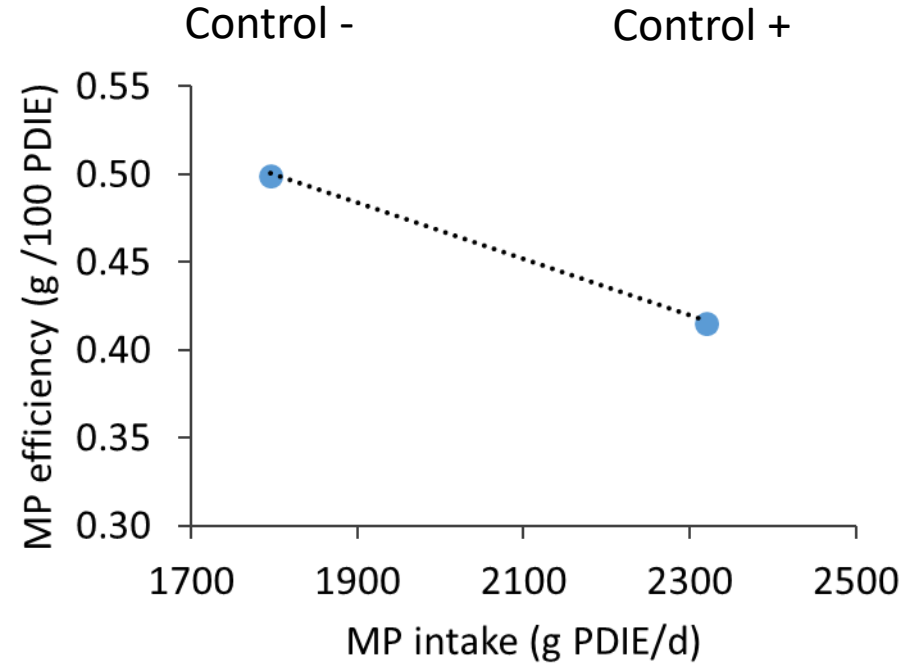
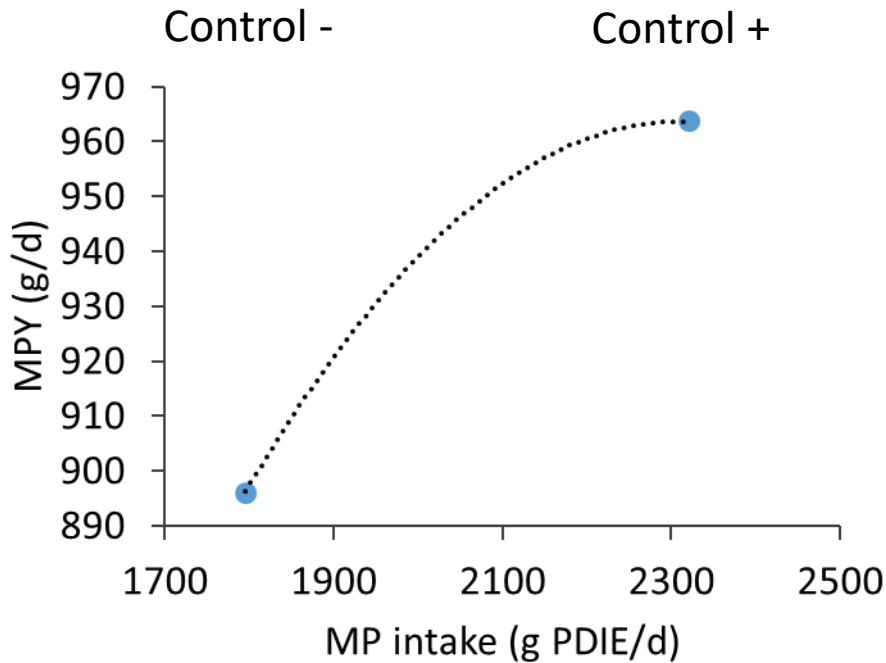
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- $MP\ eff = MPY / MP\ intake$
- Assumption of linearity

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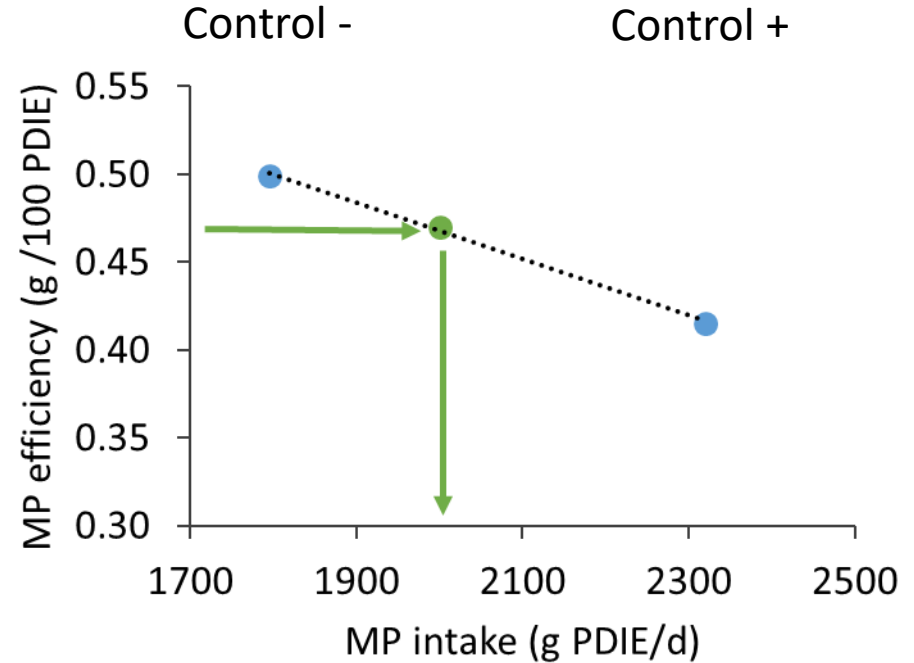
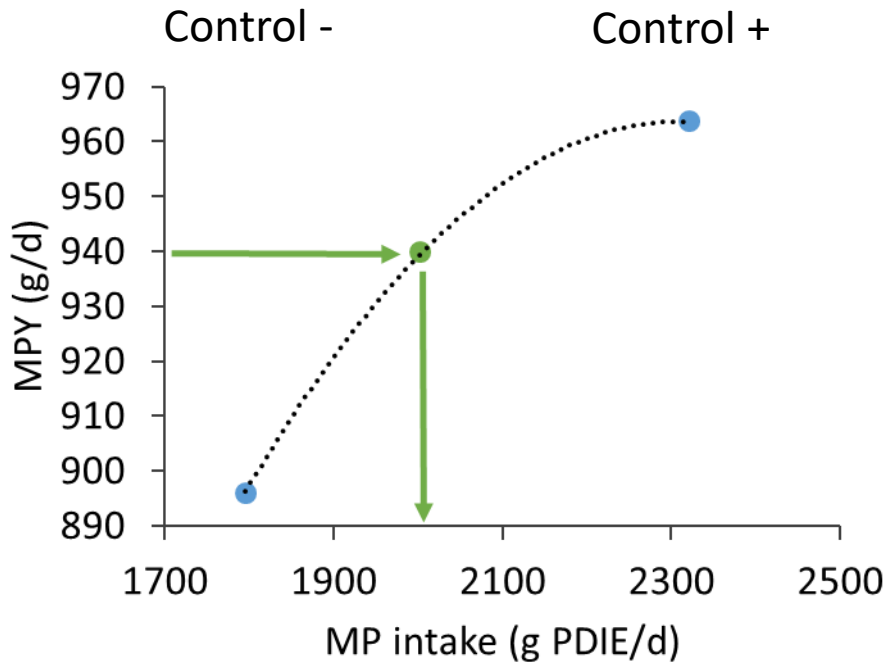
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Diets

- Same basal ration: 78% maize silage, 9.5% dehydrated lucerne, 7.5% soya bean meal, 5% energetic concentrate + mineral
- 4 dietary treatments : 2 known controls + 2 medium treatments

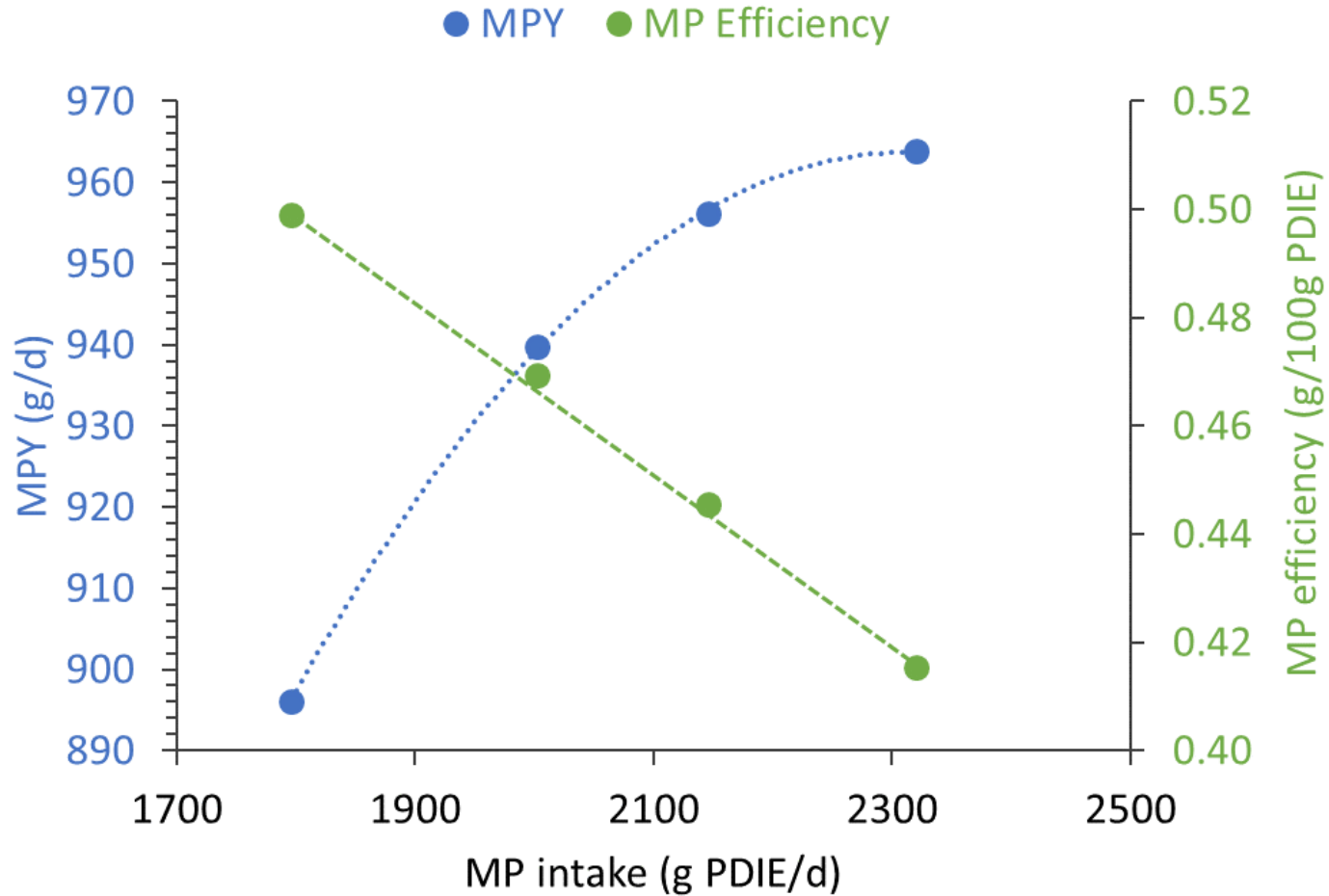
Dietary treatments	Soya bean meal (kg)	Energetic concentrate (kg)	Urea (kg)	MP (g PDIE/kg DM)
Control -	0	3.4	0.27	97
Medium Low	1.2	2.3	0.17	151
Medium high	2.2	1.0	0.09	204
Control +	3.3	0	0	252

- Avoid variations in MPY responses:
 - Energy: limited quantities
 - Microbial synthesis: same and not limiting degradable N supplies (urea)

Experimental design

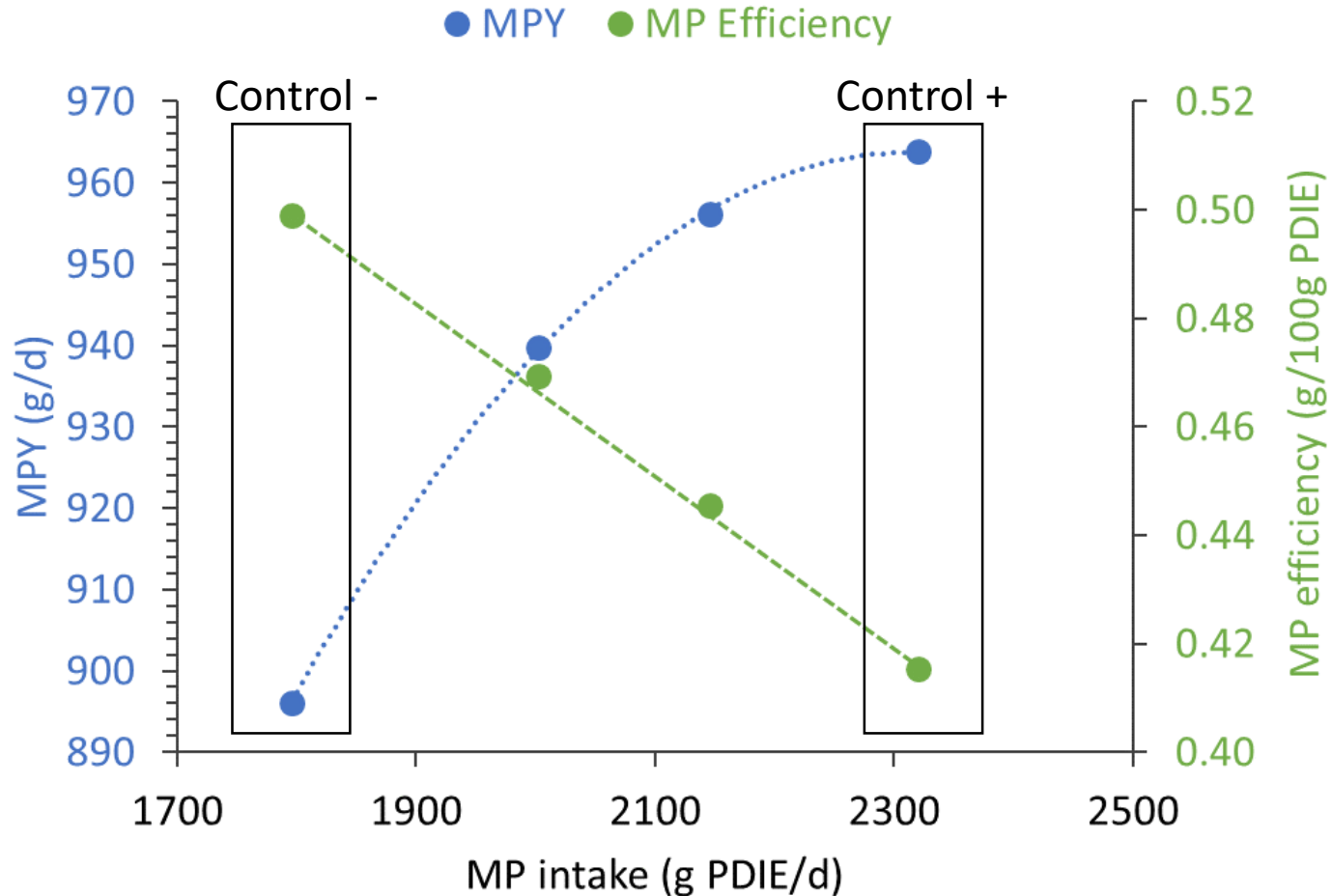
- 24 cows in 6 blocks of 4 homogenous cows
 - Repartition based on production level and parity
- 4 dietary treatments: Latin square design in each block
- 4 periods of 3 weeks
- Measurements: milk production + composition

MPY and MP efficiency response



- MPY response: 67 g/d
- MP efficiency: slightly curved

MPY and MP efficiency response



- Estimate protein value of the 2 medium treatments from their MPY response
- Comparison to theoretical values (INRA 2007)

Estimation of protein value

Dietary treatments	Theoretical value (g PDIE/kg DM)	Estimated value (g PDIE/kg DM)	Difference
Medium Low	151	165	+14
Medium High	204	221	+17

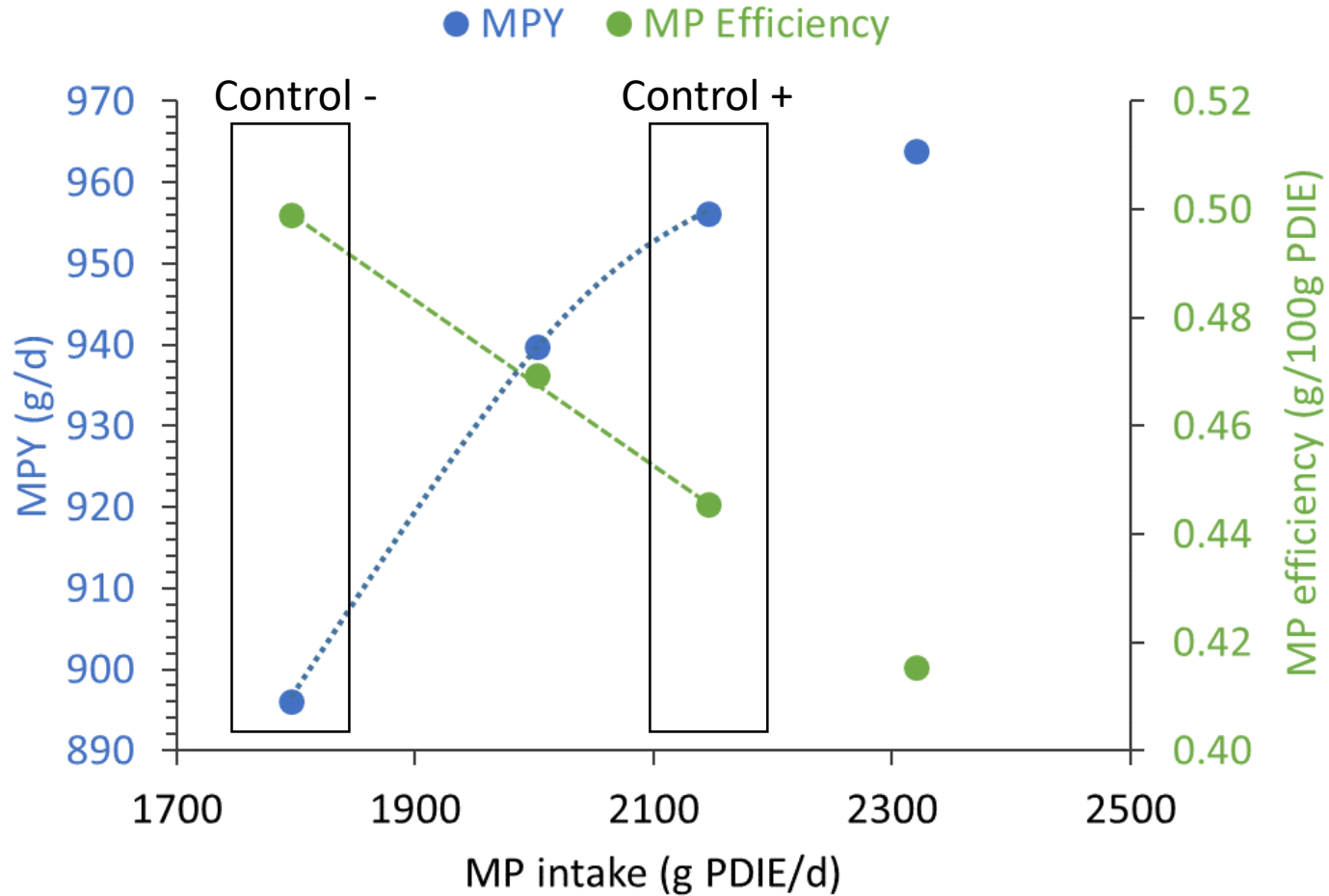
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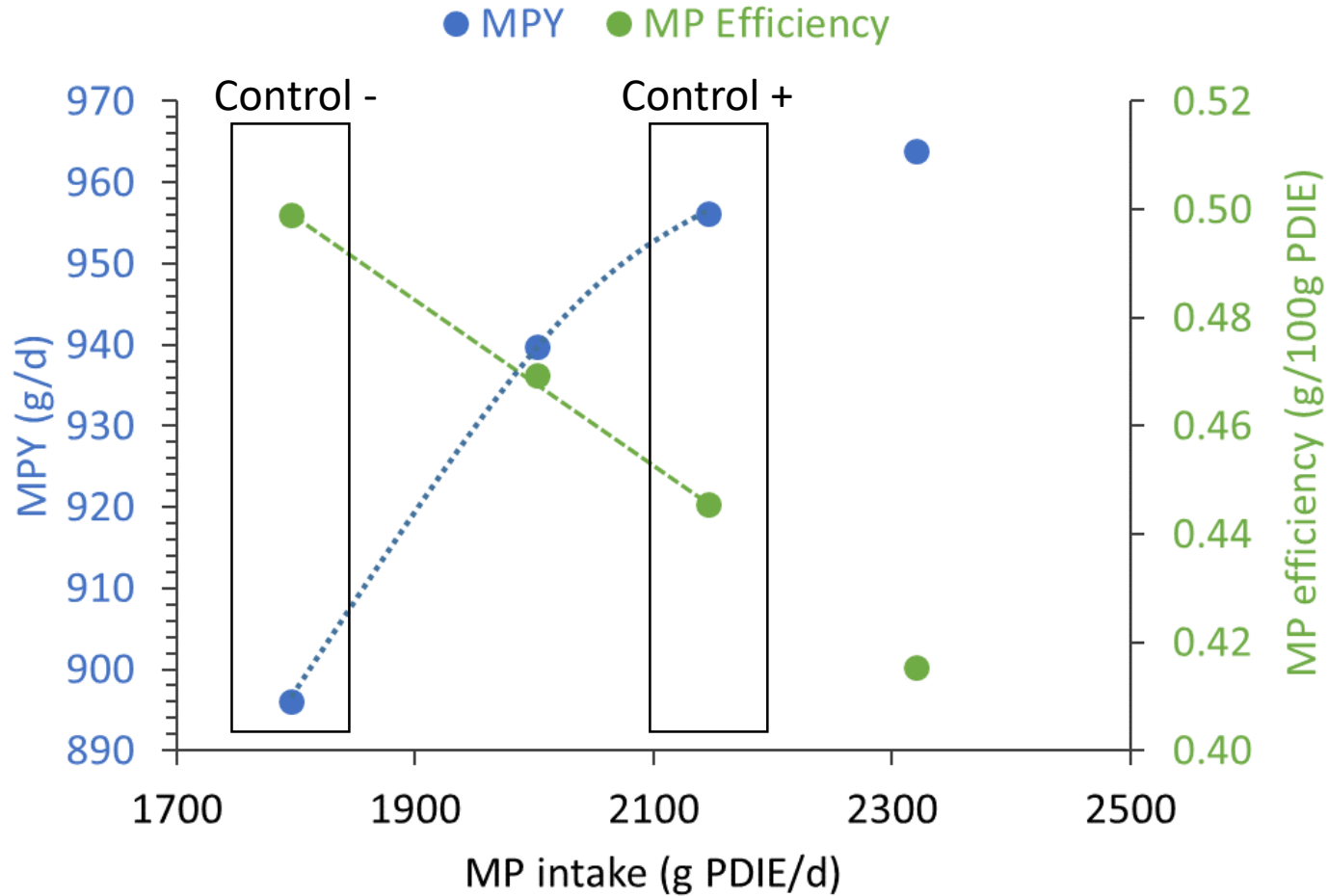
- Positive bias in the estimation
- Influence of Control + ?
 - In the upper part of MPY response = slow MPY response to protein supplies

Influence of control +



- Reduce the value of control +
- More important response in MPY

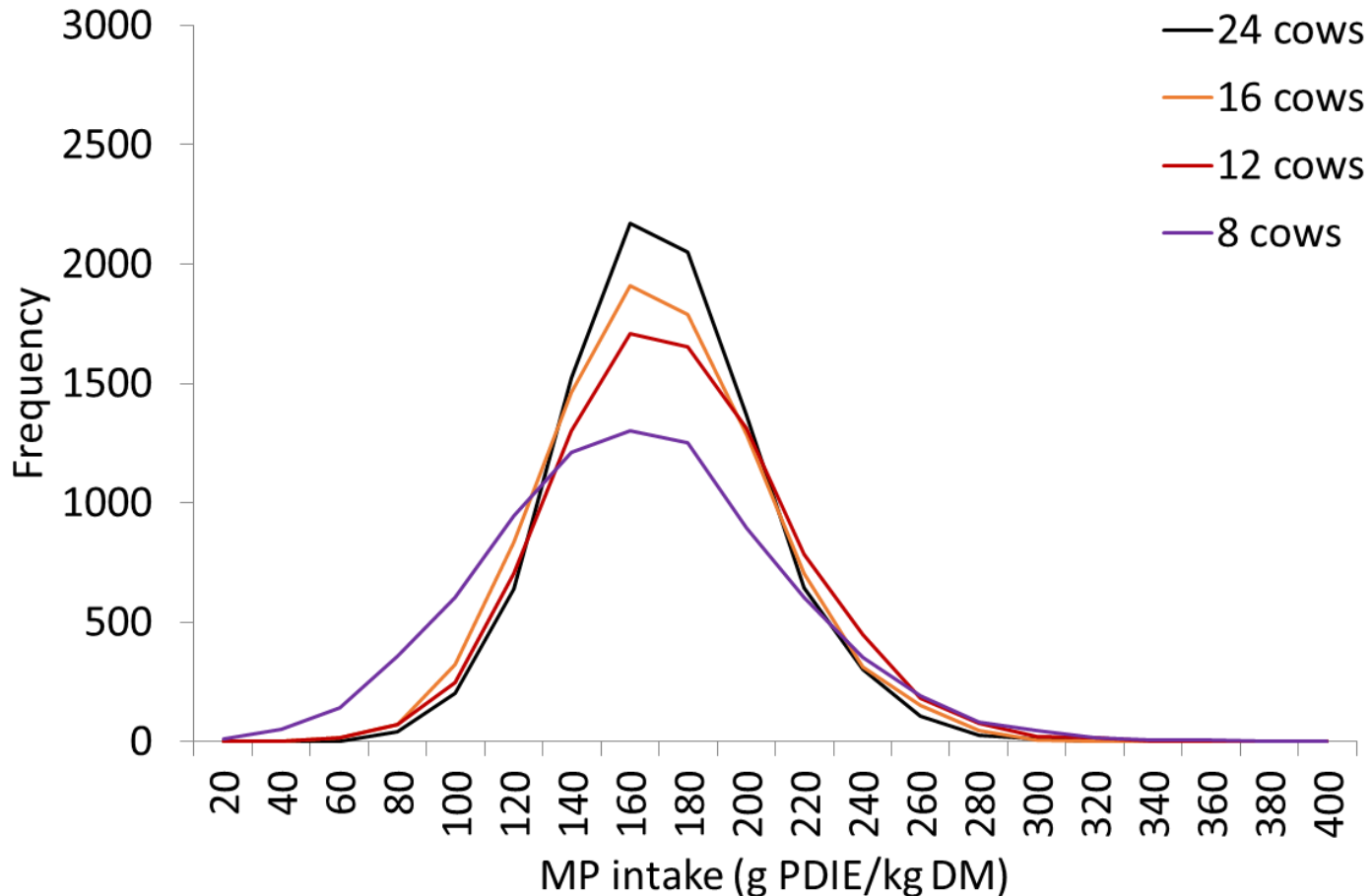
Influence of control +



Dietary treatments	Theoretical value (g PDIE/kg DM)	Estimated value (g PDIE/kg DM)	Difference
Medium Low	151	158	+7

Influence of the number of cows

- Uncertainty analysis on the Medium Low dietary treatment



- Under a number of 12 cows
 - Higher variability of estimates
 - Increase risk of bias

Conclusion

- MPY response seems to be a relevant approach to estimate feeds protein value
- Easy method, applicable in facilities equipped with feeds dispensers with no experimental measurements
- Estimation of all kind of protein protection
 - Based on physical link with feeds: heat treatment
 - Targeted microbial activity: essential oils, vegetable tannins
- Special attention in diets formulation
 - Avoid variations in MPY response (energy and microbial synthesis)
 - Avoid the range where MPY response is low (rich-protein supply)