

Evaluating the predictive ability of a mechanistic model of nitrogen partitioning applied to lactating dairy cows consuming ryegrass-based diets

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Abstract

The ability to understand and predict ruminant nitrogen (N) partitioning has become important as concerns over the environmental impact of pastoral agriculture increase. A mechanistic model of N partitioning from dairy cows consuming ryegrass was used to quantify the amount of N secreted in milk, urine, and faeces, and retained in body tissues. Data from lactating dairy cows (13 N balance experiments) were used to evaluate the model. Predictions for milk, urinary, faecal, and total excreta N were in agreement with observed values. Empirical models have been successful in predicting N partitioning from ruminants, but such approaches lack the underlying biological processes and may not apply to all observed values. An improved representation of the underlying mechanisms of N partitioning and the possibility of identifying areas where knowledge is lacking was achieved by the use of the current mechanistic model.

Keywords: Nitrogen, milk N, urinary N, faecal N, empirical, mechanistic.

Introduction

Nitrogen (N) in animal excreta, urinary N in particular, represents the largest source of N losses to the environment from pastoral agriculture. Losses of N via nitrous oxide (N₂O) and nitrate (NO₃⁻) leaching from pastoral dairying can be considerable; about 85% of agricultural N₂O emissions in New Zealand result from N excreted by grazing ruminants, and annual losses from intensively grazed pastures can be as high as 200 kg N per ha via NO₃⁻ leaching (Ledgard *et al.* 2008). Although N is an important dietary nutrient for high producing dairy cows, and adequate N in dairy diets is required to maximise production and profitability, N supply from ryegrass-based pastures is frequently in excess of requirements, whereas carbohydrate supply is often limited (Pacheco & Waghorn 2007). Consequently, the ability to understand and accurately predict the partitioning of N in ruminants becomes increasingly important.

Several mechanistic and empirical models have been developed to examine N partitioning from dairy

cows under different nutritional regimes (AFRC 1993; Castillo *et al.* 2000; Kebreab *et al.* 2002), but fewer include pasture-based diets (Pacheco *et al.* 2007). Empirical models have been successful in many situations, but this approach lacks understanding of the underlying biological processes and thus may not be applicable to conditions outside those of its development. Process-based models are more complex but offer the opportunity to reliably study the effects of N intake on N partitioning under a wide range of conditions. The objectives of this study were to (i) assess, via a dynamic model of N metabolism, the efficiency of N utilisation in dairy cows, with a focus on the amounts of N secreted (N in milk), excreted (N in urine and faeces) and retained (accretion of N in body tissues) from ryegrass-based diets, and (ii) identify gaps in the current knowledge of N partitioning within the dairy cow to focus future investigation.

Methods

Model overview and assumptions

The dynamic model of N partitioning by Kebreab *et al.* (2002) was used to quantify N flows and sinks in dairy cows consuming a ryegrass-based pasture. The model is based on a metabolisable energy (ME) and protein system (AFRC 1993), and was originally developed from diets consisting of conserved forages and concentrates. The model includes four state variables representing dietary (Di), amino acid (AA), microbial (Mi) and urea and ammonia (Ur) N (Figure 1).

A number of assumptions were made to calculate the flows between different N pools. Flows from the dietary N pool assumed rumen-undegradable protein (RUP) and rumen-degradable protein (RDP) values of 33 and 65%, respectively, and the quantity of preformed AA incorporated into microbial cells was set at 5% of potentially rumen degradable N (Kebreab *et al.* 2002). The proportion of digestible RUP N absorbed post-ruminally as AA was set at 0.79 of RUP N (AFRC 1993), and microbial true protein (75% of total microbial N) was assumed to be 85% digestible.

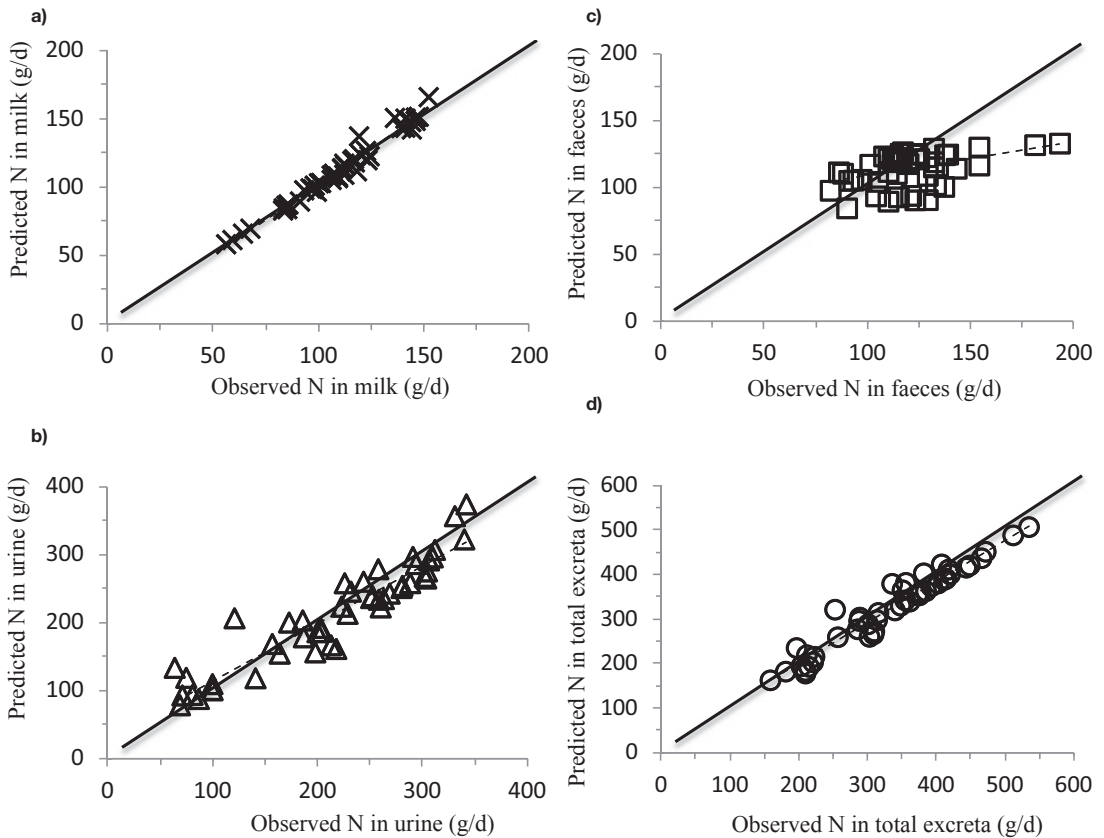


Figure 2 Relationships between observed and predicted values of N in a) milk (\times), b) urine (Δ), c) faeces (\square) and d) total excreta (\circ). Solid lines represent agreement relative to a 45° line through the origin whereas dashed lines represent the best linear fit between variables.

of N utilisation and the ratio of CP to WSC for perennial ryegrasses.

Urine and faecal excretion accounted for almost all remaining N input. As energy became limiting for MPS, the relative contribution of microbial faecal N to total faecal N diminished. Conversely, as energy supply is often rate-limiting for MPS from grass-based diets, increases in N without adequate supplies of energy resulted in surplus N, which was largely excreted via urine. In agreement with previous findings (Castillo *et al.* 2000; Kebreab *et al.* 2002), the relationship between

urine N and intake N showed an inflection point at *ca.* 450 g N/day, above which the predicted values of urinary N output were under-predicted. Unlike Castillo *et al.* (2000), where excretion of N in urine above 400 g N intake/day increased exponentially, excretion of N in urine followed a quadratic or bi-linear relationship (results not shown).

It is interesting to note that, based on the same measures of predictive ability, the N partitioning approach used by OVERSEER[®] [$N_{\text{urine}} = (11.0_{(\pm 1.1)} \times \%N \text{ in diet}) + 31.8_{(\pm 3.5)}$] produced estimates of N

Table 2 Measures of fitness (RMSPE, RPE, CCC)¹ of predicted vs. observed values of N partitioning in lactating dairy cows fed ryegrass-based diets.

Variables	RMSPE	RPE	CCC
N in milk	4.2	3.8	0.98
N in urine	25.9	12.2	0.93
N in faeces	11.5	10.1	0.36
N in total excreta	21.6	6.6	0.96

¹RMSPE is the root of the mean square prediction error (g/day); RPE is the relative prediction error (%); and CCC is the concordance correlation coefficient (scale 0 to 1).

in urine in line with the experimental data used here (RMSPE = 19.9 g/day, RPE = 9.3%, CCC = 0.95). This shows that simple empirical models can produce reasonable estimates of N excretion. However, the lack of a dynamic representation of dietary and metabolism changes makes such models less sensitive to variations in feeding strategy. Accounting for these variations is important, for instance, when analysing “what if” scenarios altering dietary strategies to address environmental issues.

Two recognised gaps in current knowledge relative to N partitioning prediction are noted. The first area relates to preformed AA uptake by microorganisms that contributes to the microbial N pool, particularly from pasture-based diets. Uptake of preformed peptides spares the cost of *de novo* microbial AA synthesis. Plant CP undergoes intensive degradation in the rumen, vastly to ammonia; nitrogenous components kinetics from ryegrasses harvested at different stages of maturity showed that the proportion of CP as RDP averaged 0.83, and maturation did not affect the ratio of RDP to RUP (Chaves *et al.* 2006). The process of intensive plant proteolysis in the rumen is suggested to be exacerbated by increased feeding frequency in grazing situations (Chen *et al.* 1987). An adequate and dynamic range of RDP/RUP values will be expected if the underlying mechanisms of reduced N loss management (i.e. timing of herbage allocation, grazing interval, the use of high sugar grasses, the use and timing of energy-dense, low protein supplements) are to be understood and implemented in grazing systems.

The second area relates to body N accretion/degradation during the entire lactation. Studies that have quantified N partition during different stages of lactation are scarce, and refer mostly to confinement-type diets (Andrew *et al.* 1994; Chilliard *et al.* 1991) or predict body lipid changes (Friggens *et al.* 2004). Although fluctuations may be minor relative to dietary N, body deposition of N may become important during the first weeks of lactation, mid to late lactation, and during the dry period. Accretion of N by body tissues of lactating dairy cows offered ryegrass-based diets accounted for up to 89 and 54 g N/day during mid lactation (>140 DIM; Miller *et al.* 2000).

Nitrogen intake continues to be the main driver of N excretion, but the level of N intake (modulated by spatial and temporal fluctuations) relative to animal requirements (modulated by milk production and body protein accretion/degradation) modifies this response. Model predictions of milk, urinary, faecal and total excreta N were in agreement with observed values. In order to improve our current understanding on how N is partitioned within the dairy cow, more research needs to be conducted in the areas of preformed amino acid

uptake by microorganisms in the rumen, particularly from pasture-based diets; and how body N accretion/degradation is affected by stage of lactation.

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