



Evaluation of between-cow variation in milk urea and rumen ammonia nitrogen concentrations and the association with nitrogen utilization and diet digestibility in lactating cows

P. Huhtanen,^{*1} E. H. Cabezas-Garcia,^{*} S. J. Krizsan,^{*} and K. J. Shingfield^{†‡}

^{*}Department of Agricultural Research for Northern Sweden, Swedish University of Agricultural Sciences, S-90183 Umeå, Sweden

[†]Natural Resources Institute Finland, Animal Production Research, FI 31600 Jokioinen, Finland

[‡]Institute of Biological, Environmental and Rural Sciences, Aberystwyth University, Aberystwyth, SY23 3EB, United Kingdom

ABSTRACT

Concentrations of milk urea N (MUN) are influenced by dietary crude protein concentration and intake and could therefore be used as a biomarker of the efficiency of N utilization for milk production (milk N/N intake; MNE) in lactating cows. In the present investigation, data from milk-production trials (production data set; n = 1,804 cow/period observations from 21 change-over studies) and metabolic studies involving measurements of nutrient flow at the omasum in lactating cows (flow data set; n = 450 cow/period observations from 29 studies) were used to evaluate the influence of between-cow variation on the relationship of MUN with MNE, urinary N (UN) output, and diet digestibility. All measurements were made on cows fed diets based on grass silage supplemented with a range of protein supplements. Data were analyzed by mixed-model regression analysis with diet within experiment and period within experiment as random effects, allowing the effect of diet and period to be excluded. Between-cow coefficient of variation in MUN concentration and MNE was 0.13 and 0.07 in the production data set and 0.11 and 0.08 in the flow data set, respectively. Based on residual variance, the best model for predicting MNE developed from the production data set was $MNE (g/kg) = 238 + 7.0 \times \text{milk yield (MY; kg/d)} - 0.064 \times MY^2 - 2.7 \times \text{MUN (mg/dL)} - 0.10 \times \text{body weight (kg)}$. For the flow data set, including both MUN and rumen ammonia N concentration with MY in the model accounted for more variation in MNE than when either term was used with MY alone. The best model for predicting UN excretion developed from the production data set (n = 443) was $UN (g/d) = -29 + 4.3 \times \text{dry matter intake (kg/d)} + 4.3 \times \text{MUN} + 0.14 \times \text{body weight}$. Between-cow variation had a smaller influence on the association of MUN with MNE and UN output than published estimates of these relationships based on treatment means, in which

differences in MUN generally arise from variation in dietary crude protein concentration. For the flow data set, between-cow variation in MUN and rumen ammonia N concentrations was positively associated with total-tract organic matter digestibility. In conclusion, evaluation of phenotypic variation in MUN indicated that between-cow variation in MUN had a smaller effect on MNE compared with published responses of MUN to dietary crude protein concentration, suggesting that a closer control over diet composition relative to requirements has greater potential to improve MNE and lower UN on farm than genetic selection.

Key words: dairy cow, digestibility, milk urea, rumen ammonia, urinary nitrogen excretion

INTRODUCTION

Dairy farming is known to cause atmospheric and hydrospheric pollution (Tamminga, 1992). Animal manure contributes to N pollution as ammonia and nitrous oxide volatilization into the atmosphere, nitrate leakage in ground water, and N runoff in surface water. Despite considerable research effort to improve the efficiency of N utilization for milk production (milk N/N intake; MNE), the conversion of dietary N into milk protein is relatively low. Over a wide range of diets, MNE averaged 247 and 277 g/kg in lactating cows fed diets typical for North America and north Europe, respectively (Huhtanen and Hristov, 2009). However, considerable variation in MNE between individual cows and across different diets highlights the potential for improving N utilization. Overfeeding of dietary CP is often the main reason for low MNE. Increased dietary CP intake has been shown to exponentially increase the proportion of N excreted in urine (Kebreab et al., 2002). Therefore, accurate determination of ruminant protein requirements and the supply of AA available for absorption are critical for optimizing animal performance, while minimizing N inputs and N emissions from milk-production systems. Under commercial conditions, monitoring the adequacy of CP in the diet requires the use of reliable diagnostic biomarkers. Urea

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¹Corresponding author: pekka.huhtanen@slu.se

in blood is the major end product of N metabolism in ruminants. Even though BUN concentration could be used as an indicator of efficient N utilization, it cannot be measured routinely on farm.

Concentrations of urea in milk and blood are closely associated in lactating cows (Broderick and Clayton, 1997). Several studies have provided evidence to support the measurement of MUN concentration in bulk-tank milk as a useful indicator of on-farm efficiency of N utilization and urinary N (UN) output (Jonker et al., 1998; Kauffman and St-Pierre, 2001; Kohn et al., 2002). Extensive evaluation of experimental data has identified dietary CP content as the major factor influencing MUN concentration (Nousiainen et al., 2004). Dietary CP content has also been reported to be a better predictor of MUN concentration (Nousiainen et al., 2004) and UN excretion (Spek et al., 2013) than estimates of protein balance in the rumen or RDP supply in excess of predicted requirements, respectively. Across a range of diets, increases in MUN concentration have been shown to be associated with a curvilinear increase in milk protein yield, reaching a nadir around 25 mg/dL, and a quadratic decrease in MNE (Nousiainen et al., 2004). Such relationships highlight that the marginal increase in milk protein yield and the associated decline in MNE to additional dietary CP can be expected to be greater for cows producing milk with a low MUN concentration. Evaluation of N intake and flow of N at the omasum measured in cows fed a range of diets indicated that zero rumen N balance was associated with an average MUN concentration of 8.3 mg/dL (Broderick et al., 2010), with the implication that a MUN below this concentration might reflect a deficiency in RDP.

Most assessments of the potential of measuring MUN concentrations as a diagnostic of MNE and biomarker of N metabolism and UN output in lactating cows have been based on treatment means (Jonker et al., 2002; Nousiainen et al., 2004; Spek et al., 2013). However, a recent study reported differences in MUN concentration that were not explained by variations in dietary CP intake, milk yield, or BW (Aguilar et al., 2012), with the corollary that measurements of MUN may not be a robust biomarker for individual cows or a useful phenotyping tool in genetic selection for improved MNE or lower UN output. Even though variation in MUN concentration due to cow is widely recognized (Wattiaux et al., 2005; Stoop et al., 2007; König et al., 2008), the influence of between-cow variation on the association of MUN concentrations with MNE, UN, or overall N metabolism is not well defined. The objective of this study was to characterize the extent of between-animal variation in the concentrations of MUN or rumen ammonia nitrogen (RAN), and the influence

of between-cow variation on the association of these parameters with MNE, UN excretion, and total-tract diet digestibility. Evaluations were made on data from milk-production trials and metabolic studies in lactating cows using mixed-model regression analysis that allowed the effects of diet and period to be removed.

MATERIALS AND METHODS

Experimental Data

Evaluations were performed using 2 data sets, one based on measurements from cows used in milk-production trials and the other containing a smaller set of more detailed measurements from metabolic studies in lactating cows. The production data set (Supplemental Table S1, <http://dx.doi.org/10.3168/jds.2014-8215>) comprised individual cow/period observations ($n = 1,978$) obtained from 21 change-over milk-production trials. Data originated from 4 production trials conducted in Sweden, 2 in Norway, and 15 in Finland. All production trials were conducted as either Latin square or cyclic change-over designs. In 17 trials, forages were fed ad libitum and concentrates were offered on a flat-rate basis, whereas in the rest a TMR was offered ad libitum. Grass silage was the main forage component for all diets, but in 6 trials, grass silage was partially replaced with red clover or whole-crop cereal silages. Concentrate supplements contained oats and barley, fibrous by-products from the food industry, and protein supplements, typically soybean meal, and rapeseed meal or rapeseed expeller. All rapeseed feeds were of double-zero varieties with low glucosinolate content. Measurements included parity (primiparous or multiparous), DIM, BW, feed intake, diet chemical composition, including silage fermentation characteristics, milk yield, milk composition, and MUN concentration. Determinations of diet chemical composition and measurements of DMI were used to calculate the supplies of ME and MP using the Finnish feed evaluation system (MTT, 2013). The yield of ECM was calculated according to Sjaunja et al. (1991). For 9 milk-production trials MUN concentrations were determined as ammonia following urease hydrolysis (McCullough, 1967) in a single laboratory. Concentrations of MUN in the other 12 milk-production trials was measured by infrared spectroscopy (MilkoScan 6000, Foss Electric, Hillerød, Denmark) in 3 different laboratories. Observations ($n = 174$) for cows <20 DIM, producing <10 kg of milk/d, with a calculated ME balance of >+50 MJ/d or <-50 MJ/d were excluded from the production data set.

Total-tract diet digestibility ($n = 443$) was determined in 10 milk-production trials. Measurements of diet di-

gestibility for individual cows were made on average for 50% of the animals recruited to a milk-production trial based on twice-daily collection of spot fecal samples over 5 d using acid insoluble ash as an internal marker (Van Keulen and Young, 1977). Urinary N excretion was not measured but calculated as

$$\text{UN} = \text{N intake} - \text{fecal N} - \text{milk N} - \text{retained N.}$$

Retained N was estimated from calculated ME balance assuming that for cows in positive energy balance, 1.0-kg-of-BW change was equivalent to 34 MJ of ME, whereas 1.0-kg-of-BW change during periods of negative energy balance was equivalent to 28 MJ (MTT, 2013). Each kilogram of BW change was assumed to contain 160 g of CP (MTT, 2013).

Data from metabolic studies in lactating cows (flow data set) included measurements of nutrient intake, rumen fermentation characteristics, nutrient flow at the omasum, milk yield, and milk composition (Supplemental Table S2, <http://dx.doi.org/10.3168/jds.2014-8215>). The flow data set comprised 450 cow/period observations from 29 experiments. Total-tract diet digestibility was determined by total fecal collection in 23 experiments or from the collection of spot fecal samples using acid insoluble ash as a marker in 6 experiments. Measurements of nutrient flow at the omasum were made in 22 studies ($n = 331$) using the omasal sampling technique and a triple-marker system. Concentrations of RAN were determined in all 29 experiments ($n = 450$). In 20 studies, MUN concentration was measured by urease hydrolysis (McCullough, 1967) in a single laboratory. For all 29 metabolic studies, cows were offered diets of similar composition to that fed in the milk-production trials.

Statistical Analysis

Estimates of variance components were evaluated using the PROC MIXED procedure of SAS (version 9.3; SAS Institute Inc., Cary, NC) with experiment (**Exp**), diet within experiment [**Diet(Exp)**], period within experiment [**Period(Exp)**], and cow within experiment [**Cow(Exp)**] as random factors. Covariance structure was specified using the TYPE = VC option in the RANDOM statement. The standard deviation and coefficient of variation for each factor (Tables 3 and 4) were calculated as the square root of the variance estimate and standard deviation divided by the respective mean value of each factor (Tables 1 and 2), respectively.

Repeatability values (Rep) for MUN and RAN concentrations were calculated as

$$\text{Rep} = \sigma_{\text{Cow}}^2 / (\sigma_{\text{Cow}}^2 + \sigma_{\text{Residual}}^2),$$

where σ_{Cow}^2 and $\sigma_{\text{Residual}}^2$ are Cow(Exp) and residual variances, respectively. Repeatability values provide an estimate of the correlation between values from consecutive samples on the same cow, on the same diet, and within the same period of the same experiment.

To explore the relationships between the variables of interest [MUN, MNE, UN, total-tract OM digestibility (**OMD**), and total-tract NDF digestibility (**NDFD**)], data were analyzed by regression analysis within the MIXED procedure of SAS (Littell et al., 2006) using the following model:

$$Y_{ij} = B_0 + B_1X_{1ij} + b_0 + b_1X_{1ij} + B_2X_{2ij} + B_3X_{3ij} + B_4X_{4ij} + e_{ij},$$

where Y_{ij} = the expected value for the dependent variable Y observed at level of j of the independent variable X in the study i ; B_0 = the overall intercept (fixed effect); b_0 = the random effect of study i on the intercept ($i = 1, \dots, 21$ for the production data set; $i = 1, \dots, 29$ for the flow data set); B_1, \dots, B_4 are regression coefficients of Y on X_1, \dots, X_4 of Y across all studies (fixed effects); X_{1ij}, \dots, X_{4ij} = value j of the continuous variable X_1, \dots, X_4 in study i ; b_i = the random effect of study i on the regression coefficient of Y on X_1 in study i ($i = 1, \dots, 21$ for the production data set; $i = 1, \dots, 29$ for the flow data set); and e_{ij} = the residual error.

In different models, the number of independent variables varied between 1 and 4. The model included 2 random statements: a random intercept and slope of X_1 with SUBJECT = Diet(Exp), and a random intercept with SUBJECT = Period(Exp), using the TYPE = VC (variance components) covariance structure for both random statements. The method = ML (maximum likelihood) statement was used in the PROC MIXED model syntax. Only one random independent variable was used to avoid overparameterized models and improve convergence (St-Pierre, 2001). The models were evaluated on the basis of Akaike's information criterion with correction (**AICc**) and residual variance. Syntaxes of statistical programs are available in Supplemental Table S3 (<http://dx.doi.org/10.3168/jds.2014-8215>).

RESULTS

Evaluation Data Sets

Mean diet characteristics and animal variables for the production data set are shown in Table 1. Both animal and diet parameters exhibited considerable variation and covered the range in dietary chemical composition relevant to commercial milk production. Variation in

Table 1. Description of diet composition, intake, and milk yield and composition for the production data set derived from 21 milk-production trials (n = 1,804)

Item	Mean	SD	Minimum	Maximum
Intake, kg/d				
Forage DM	12.5	2.35	4.1	21.0
Concentrate DM	8.3	2.39	2.5	14.6
Total DM	20.8	3.04	10.9	30.2
ME intake, MJ/d	236	36.5	113	352
ME balance, MJ/d	14	17.5	-48	50
MP supply, kg/d	1.96	0.324	0.87	3.02
Diet composition				
OM, g/kg of DM	925	9.7	887	948
CP, g/kg of DM	162	19.7	111	220
NDF, g/kg of DM	437	41.6	313	563
ME, MJ/kg of DM	11.3	0.44	10.2	12.4
Milk yield and composition				
Milk, kg/d	29.7	6.42	10.1	51.1
ECM, kg/d	30.9	5.88	12.2	52.3
Fat, g/kg	43.7	6.23	27.3	66.6
Protein, g/kg	33.7	3.16	25.4	46.8
Lactose, g/kg	47.7	2.19	38.3	54.8
MUN, mg/dL	12.0	3.52	2.2	28.8
MNE, ¹ g/kg	296	39.7	176	444
Feed efficiency, kg of ECM/kg of DMI	1.48	0.158	0.91	2.22
BW, kg	609	68.8	443	822
DIM	131	51.6	23	349

¹Efficiency of N utilization for milk production defined as milk N/N intake.

these parameters was similar in the flow data set (Table 2). For both data sets, the range in diet composition was similar, whereas average DMI and milk yield were 1.4 and 2.6 kg/d lower in the production compared with the flow data set, respectively. Concentrations of MUN for individual cows varied substantially (CV = 0.286 and 0.317 in the production and flow data sets, respectively).

Variance estimates of dependent variables for the production and flow data sets are presented in Tables 3 and 4, respectively. Differences between experiments was the largest source of variation in the production data set (Table 3), reflecting the differences in diet composition between experiments and variance introduced by differences in analytical methods. Least squares means of MUN concentration using CP and ME intakes as covariates were 9.5, 8.9, and 13.1 mg/dL for the 3 participating laboratories using infrared reflectance spectroscopy (n = 240, 124, and 705, respectively), and 12.0 mg/dL for the laboratory using urease hydrolysis (n = 735).

For MUN concentration, the extent of between-cow variability and variation due to Diet(Exp) for the production data set were of a similar magnitude (CV = 0.132 and 0.125, respectively; Table 3). Repeatability of MUN concentration (0.66) in the production data set was substantially lower than for milk fat (0.76) or milk protein (0.91) content.

Estimates of variance components for MNE were similar in the production and flow data sets (Tables

3 and 4, respectively), with the exception of greater between-study variability in the flow data set (0.122 vs. 0.076). For both data sets, variation in MNE was smaller than for MUN concentration (Tables 1 and 2).

Concentration of RAN in the flow data set was more variable (CV = 0.515; Table 2) than MUN concentration in both the production and flow data sets (Tables 1 and 2). The variance component of RAN concentration (Table 4) associated with Cow(Exp) was highly significant ($P < 0.001$) and of greater magnitude than for MUN concentration in the same data set (CV = 0.144 vs. 0.106). Total variance of MNE and its distribution were similar in both data sets (Tables 3 and 4). Experiment and Diet(Exp) were the largest sources of variation in calculated UN output (Table 3). Between-cow variability in OMD was rather small (CV = 0.012 and 0.016 in the production and flow data sets, respectively).

Nitrogen Intake and Excretion

Distribution of N intake in the production data set is shown in Table 5. The proportion of N intake lost in urine exhibited more variation compared with other routes of N excretion (Table 5). Partitioning of N intake in milk was lower than for feces. Regression based on the mixed model analysis of milk N, fecal N, and urine N output or retained N against N intake indicated that incremental increases in N intake were partitioned in milk (0.262 ± 0.005), feces (0.272 ± 0.012), urine (0.374

Table 2. Description of diet composition, intake, milk production, total-tract nutrient digestibility, and N flow at the omasum for the flow data set derived from 29 metabolic studies in lactating cows

Item	n	Mean	SD	Minimum	Maximum
Intake, kg/d					
Forage DM	450	11.8	2.42	2.5	18.8
Concentrate DM	450	7.7	2.46	0.0	17.3
Total DM	450	19.5	2.73	11.4	27.5
Diet composition					
CP, g/kg of DM	450	159	21.0	126	239
NDF, g/kg of DM	450	401	55.6	288	534
Milk yield and composition					
Milk, kg/d	450	27.2	6.99	5.0	44.6
ECM, kg/d	450	27.2	6.24	5.2	47.4
Fat, g/kg	450	40.6	6.04	16.5	55.9
Protein, g/kg	450	33.4	3.97	21.3	58.0
Lactose, g/kg	450	47.8	2.20	37.1	53.7
MUN, mg/dL	309	10.2	3.24	3.3	21.1
MNE, ¹ g/kg	450	291	51.9	95	475
BW, kg	295	624	64.5	459	789
Total-tract digestibility, g/kg					
OM	414	739	34.4	641	857
CP	414	683	51.5	519	876
NDF	414	648	69.2	447	828
Flow at the omasum					
NAN, g/d	331	474	100.8	226	667
Microbial N, g/d	331	311	72.4	112	496
Microbial N, g/kg of DOMR ²	295	24.5	3.81	15.4	37.0
RAN, ³ mg/dL	450	9.7	5.01	2.0	34.0

¹Efficiency of N utilization for milk production defined as milk N/N intake.

²Organic matter apparently digested in the rumen.

³Rumen ammonia N concentration.

± 0.018), and body tissues (0.056 ± 0.004). Mean estimated N retention in the production data set was close to zero (+8 g/d; Table 5).

For the 10 milk-production trials where total-tract digestibility was measured ($n = 443$), regression of apparent digestible N against dietary N concentration basis estimated true N digestibility (slope) and metabolic fecal N excretion (intercept) as 0.911 ± 0.040 and -5.8 ± 1.10 g/kg of DMI, respectively.

Factors Influencing MUN Concentration

Including Diet(Exp) and Period(Exp) effects in the statistical analysis allowed for an evaluation of animal factors influencing MUN in lactating cows independent of dietary treatment or experimental period. For the production data set, MUN concentration increased ($P < 0.001$) with increases in the yields of milk and ECM (Table 6). Furthermore, MUN concentration was associated negatively ($P < 0.001$) with milk protein concentration, whereas no relation existed ($P > 0.05$) between the concentrations of MUN and milk fat concentration. Increases in DIM had a quadratic effect ($P < 0.01$) on MUN concentration, reaching a maximum on d 123 postpartum. In a bivariate model, MUN concentration was positively associated with milk yield and negatively with milk protein concentration. Calculated ME bal-

ance had no influence ($P > 0.05$) on MUN concentration (data not presented).

Factors Influencing Efficiency of Nitrogen Utilization

In the production data set, MNE was positively associated ($P < 0.001$) with milk yield and negatively related ($P < 0.001$) with MUN concentration (Table 7). The effects of these factors were additive. Based on residual variance, statistical models resulted in a better fit when both milk yield and MUN concentration were included as independent variables. Both DIM and BW had a negative influence on MNE. The model with the best fit included milk yield, MUN concentration, and BW as independent variables. In the flow data set, MNE was positively associated with milk yield and negatively related to MUN concentration (Table 8). Analysis of the flow data set for experiments ($n = 19$) in which both RAN and MUN were analyzed ($n = 309$) indicated that the concentration of RAN resulted in a slightly better prediction of MNE compared with MUN concentration. However, based on the AICc for each model, the prediction of MNE was improved when the concentrations of both MUN and RAN were included in the model as independent variables, particularly when milk yield was also included in the model (Table 8).

Table 3. Variance-component estimates of MUN concentration, efficiency of N utilization for milk production (MNE), urinary N (UN) output, and total-tract OM digestibility (OMD) for lactating cows, developed using the production data set derived from 21 milk-production trials

Item ¹	Estimate	SE	Pr > Z ²	SD ³	CV ⁴
MUN, mg/dL, n = 1,804					
Exp	5.41	1.909	0.002	2.325	0.194
Period(Exp)	0.80	0.159	<0.001	0.893	0.075
Diet(Exp)	2.23	0.282	<0.001	1.493	0.125
Cow(Exp)	2.49	0.189	<0.001	1.579	0.132
Residual	1.28	0.056	<0.001	1.133	0.095
MNE, g/kg, n = 1,804					
Exp	502	197.9	0.006	22.4	0.076
Period(Exp)	182	36.5	<0.001	13.5	0.046
Diet(Exp)	367	46.2	<0.001	19.2	0.065
Cow(Exp)	382	29.4	<0.001	19.5	0.066
Residual	217	9.5	<0.001	14.7	0.050
UN output, g/d, n = 443					
Exp	1,594	826.2	0.027	39.9	0.206
Period(Exp)	60	22.1	0.003	7.8	0.040
Diet(Exp)	965	159.0	<0.001	31.1	0.161
Cow(Exp)	260	47.2	<0.001	16.1	0.083
Residual	262	25.2	<0.001	16.2	0.084
OMD, g/kg, n = 443					
Exp	444	233.0	0.028	21.1	0.030
Period(Exp)	87	27.7	0.001	9.3	0.013
Diet(Exp)	213	41.3	<0.001	14.6	0.020
Cow(Exp)	77	19.8	<0.001	8.8	0.012
Residual	203	19.6	<0.001	14.3	0.020

¹Exp = experiment; Period(Exp) = period within experiment; Diet(Exp) = diet within experiment; Cow(Exp) = cow within experiment.

²Probability of Z-value.

³Calculated as the square root of the variance-component estimate.

⁴Calculated as SD divided by the respective mean value of the variable.

Factors Influencing UN Excretion

In the production data set, total DMI was the best single predictor of UN excretion based on residual variance and the AICc (Table 9). Output of UN was positively ($P < 0.001$) related to MUN concentration. The regression coefficient remained rather constant (4.3 to 5.8) when MUN was used as the sole independent variable or when included in the model containing other animal variables. The regression coefficients for MUN remained similar when milk yield was used as the only independent variable or when both milk yield and MUN concentration were used to predict UN output (2.8 vs. 2.7 g/d per kg/d, respectively). Predictions of UN excretion were improved further by including DIM or BW as the third independent variable.

Factors Influencing Feed Efficiency and Total-Tract Digestibility

The influence of MUN concentration on feed efficiency was 0.010 ± 0.0036 kg of ECM/kg of DMI per unit change in MUN (mg/dL; $P < 0.01$) when milk yield and MUN concentration were used as independent variables. In the flow data set, the influence of

MUN or RAN concentration was positively associated with improvements in total-tract OMD (Table 10).

The coefficients for MUN and RAN remained unchanged when total DMI was included in the model (Table 10). Quantitatively, the effect of RAN concentration on NDFD was greater than on OMD (3.9 vs. 1.4 g/kg per mg/dL). A positive relationship also existed between MUN concentration and NDFD (model not reported). No relationship ($P > 0.10$) was found between RAN concentration and the digestibility of neutral detergent solubles (models not shown). Furthermore, the concentration of MUN or RAN had no influence ($P > 0.10$) on the flows of NAN or microbial N at the omasum (models and regression equations not presented).

DISCUSSION

Concentrations of MUN are influenced by dietary CP concentration and intake, are closely correlated with UN excretion, and can be measured at a low cost, and are therefore attractive as a biomarker of protein intake relative to requirements in lactating dairy cows (Nousiainen et al., 2004). However, the value of MUN concentration as a diagnostic of protein feeding for an individual cow or as a suitable phenotype for genetic

Table 4. Variance-component estimates of MUN concentration, efficiency of N utilization for milk production (MNE), rumen ammonia N concentration (RAN), and total-tract OM digestibility (OMD) developed using the flow data set derived from 29 metabolic studies in lactating cows

Item ¹	Estimate	SE	Pr > Z ²	SD ³	CV ⁴
MUN, mg/dL, n = 309					
Exp	3.86	1.725	0.013	1.964	0.193
Period(Exp)	1.08	0.277	<0.001	1.038	0.102
Diet(Exp)	2.83	0.619	<0.001	1.683	0.165
Cow(Exp)	1.19	0.281	<0.001	1.092	0.107
Residual	1.26	0.160	<0.001	1.125	0.110
MNE, g/kg, n = 450					
Exp	1156	383.3	0.001	34.0	0.117
Period(Exp)	221	50.7	<0.001	14.9	0.051
Diet(Exp)	317	66.2	<0.001	17.8	0.061
Cow(Exp)	505	89.4	<0.001	22.5	0.077
Residual	361	38.4	<0.001	19.0	0.065
RAN, mg/dL, n = 450					
Exp	14.28	4.310	0.001	3.779	0.389
Period(Exp)	1.81	0.396	<0.001	1.346	0.138
Diet(Exp)	2.29	0.482	<0.001	1.514	0.156
Cow(Exp)	1.91	0.399	<0.001	1.382	0.142
Residual	2.82	0.294	<0.001	1.679	0.173
OMD, g/kg, n = 414					
Exp	491	180.2	0.003	22.2	0.030
Period(Exp)	112	28.0	<0.001	10.6	0.014
Diet(Exp)	229	47.7	<0.001	15.1	0.020
Cow(Exp)	132	31.0	<0.001	11.5	0.016
Residual	214	23.6	<0.001	14.6	0.020

¹Exp = experiment; Period(Exp) = period within experiment; Diet(Exp) = diet within experiment; Cow(Exp) = cow within experiment.

²Probability of Z-value.

³Calculated as the square root of the variance-component estimate.

⁴Calculated as SD divided by the respective mean value of the variable.

selection for improved MNE or lower UN excretion is dependent on the extent of between-animal variation that cannot be explained by diet, level of production, or stage of lactation (Aguilar et al., 2012). Variability in MUN concentration (CV = 0.30) in the production and flow data sets was similar to that reported for measurements made on commercial farms (Wattiaux et al., 2005; Stoop et al., 2007; König et al., 2008). However, the repeatability of MUN concentration (0.66) was higher than the corresponding values of 0.43

reported previously (Stoop et al., 2007). It is probable that this difference is related to a greater variability in dietary CP concentration on commercial farms compared with diets fed in controlled experiments. Dietary CP concentration is the most important dietary factor affecting MUN concentration (Broderick and Clayton, 1997; Nousiainen et al., 2004). In practice, diet composition can vary between test-days, thereby lowering the repeatability of MUN determination. For experimental data used in this evaluation, some variation

Table 5. Description of the intake, excretion, and retention of N, and total-tract nutrient digestibility for the production data set derived from 21 milk-production trials (production data set, n = 443)

Item	Mean	SD	Minimum	Maximum
N, g/d				
Intake	531	104.2	259	767
Feces	172	38.7	68	266
Milk	158	32.6	62	247
Urine	193	52.1	48	379
Retained ¹	8	12.2	-40	37
Total-tract digestibility, g/kg				
OM	713	31.9	607	827
NDF	614	65.0	404	790
CP	676	42.7	525	854

¹Estimated from calculated ME balance assuming a CP concentration of 160 g/kg of BW change, 1.0 kg of BW gain is equivalent to 34 MJ of ME, and 1.0 kg of BW loss is equivalent to 28 MJ of ME (MTT, 2013).

Table 6. Influence of animal factors on MUN concentration (mg/dL) in lactating cows estimated by mixed-model regression analysis ($MUN = A + BX_1 + CX_2$) of the production data set derived from 21 milk-production trials ($n = 1,804$)¹

X_1	X_2	A^2	SE	B	SE	P -value	C^2	SE	Residual	AICc ³
MY		9.8	0.39	0.069	0.011	<0.001			3.50	7,984
ECM		9.8	0.42	0.065	0.012	<0.001			3.54	7,997
FC		12.5	0.50	-0.016	0.010	0.11			3.65	8,036
PC		16.6	0.62	-0.141	0.017	<0.001			3.50	7,966
BW		12.4	0.54	-0.0009	0.0008	0.25			3.65	8,036
DIM	DIM × DIM	10.9 ⁴	0.44	1.69	0.470	<0.001	-0.662	0.147	3.60	8,013
MY	PC	14.8	0.92	0.030	0.013	0.02	-0.115	0.019	3.41	7,951

¹MY = milk yield, kg/d; ECM, kg/d; FC = milk fat concentration, g/kg; PC = milk protein concentration, g/kg; BW is in kg.

²All P -values <0.001.

³AICc = Akaike's information criterion with correction.

⁴Coefficients calculated for DIM/100.

in dietary CP concentration may exist between cows fed the same diet, because in most of the production trials, concentrates were fed on a flat-rate basis and forages were offered ad libitum. Consequently, dietary CP concentration may fluctuate between individual cows if the CP concentration of concentrates and forages differ. For the production data set, variation due to cow within experiment explained only 0.1% of total variance in dietary CP concentration and, therefore, is not a major factor contributing to between-animal variation in MUN concentration ($CV = 0.13$; Table 3). Furthermore, between-cow and residual variance did not change when dietary CP concentration was included in the model predicting MUN concentration, which also suggests that the difference in dietary CP concentration was a small contributor to phenotypic variation in MUN concentration.

Influence of Animal Factors on Milk Urea Concentration

Use of a mixed model in the evaluation of production and flow data sets allowed the effects of diet within study and period within study to be removed, and therefore, only between-animal differences in N digestion and metabolism contributed to the relationship between milk yield and MUN concentration. Reports on the strength of the relationship between milk yield and MUN concentration based on measurements of individual animals have been inconsistent. In some studies, a negative association between MUN concentration and milk yield has been reported (Broderick and Clayton, 1997; Stoop et al., 2007). In contrast, increases in FCM yield of multiparous cows up to 58 kg/d were accompanied by an increase in MUN concentration (Wattiaux et al., 2005). A positive (0.13) phenotypic correlation between MUN and milk yield has also been reported (König et al., 2008). However, a close positive association between MUN concentration and milk yield observed in com-

mercial herds may simply reflect higher-yielding cows consuming diets containing more CP. In the evaluation of the production data set, the effect of milk yield on MUN concentration, although significant, was quantitatively small (i.e., 0.069 mg/dL per unit kg/d; Table 6). This was exemplified by a single standard deviation difference in milk yield between cows within Diet(Exp) and Period(Exp) (i.e., 4.25 kg/d or 14.3% of average milk yield; Table 1) corresponding to a $\pm 2.5\%$ difference in MUN concentration (i.e., $0.069 \times 4.25/12.0\%$; Table 1). Using the relationship between dietary CP and MUN concentrations based on a meta-analysis of treatment means ($MUN, \text{mg/dL} = -14.2 + 0.17 \times \text{dietary CP concentration, g/kg of DM}$; Nousiainen et al., 2004), and the relationship between milk yield and MUN concentration for the production data set ($MUN, \text{mg/dL} = 9.8 + 0.069 \times \text{milk yield, kg/d}$; Table 6), then the changes in MUN concentration accompanying a 1-kg increase in milk yield can be calculated to correspond to a 0.4 g/kg increase in dietary CP concentration. Such comparisons suggest that a 1-kg increase in milk yield would require an additional 8 g of dietary CP for gluconeogenesis in lactating cows with a DMI of 20 kg, assuming that deamination and use of AA carbon for glucose synthesis have the same influence on MUN concentration as an increase in CP intake. The positive effect of milk yield on MUN concentration and negative effect of milk protein concentration are in line with this suggestion.

A negative association between MUN and milk protein concentration is consistent with previous evaluations based on treatment means (Nousiainen et al., 2004). However, other studies have reported positive phenotypic and genotypic associations between MUN and milk protein concentration (Stoop et al., 2007). The reasons for a negative relationship between the concentration of MUN and milk protein are not obvious, particularly when milk protein concentration was not associated with MNE (Table 7) or UN output

Table 7. Influence of animal factors on the efficiency of N utilization for milk production (MNE; milk N/N intake, g/kg) in lactating cows estimated by mixed-model regression analysis ($MNE = A + BX_1 + CX_2 + DX_3 + EX_4$) of the production data set derived from 21 milk-production trials ($n = 1,804$)¹

X_1	X_2	X_3	X_4	A^2	SE	B^2	SE	C^2	SE	D^2	SE	E^2	SE	Residual	AICc ³
MY				218	4.2	2.61	0.115							441	16,703
MUN				313	4.3	-1.57	0.291							587	17,124
MY	MUN			242	4.7	2.77	0.115	-2.40	0.256					435	16,625
MY	MY × MY			155	11.3	6.91	0.729	-0.070	0.0118					430	16,668
MY	MY × MY	MUN		173	11.1	7.64	0.720	-0.080	0.0116	-2.54	0.254			423	16,579
MY	MY × MY	BW		218	12.3	6.24	0.706	-0.054	0.0115	-0.095	0.0084			399	16,547
MY	MY × MY	DIM		169	11.8	6.70	0.728	-0.070	0.0117	-0.059	0.0156			426	16,655
MY	MY × MY	MUN	BW	238	12.0	7.00	0.694	-0.064	0.0112	-2.65	0.245	-0.100	0.0083	388	16,442

¹MY = milk yield, kg/d; MUN is in mg/dL; BW is in kg.²All P -values ≤ 0.001 .³AICc = Akaike's information criterion with correction.**Table 8.** Influence of animal factors on the efficiency of N utilization for milk production (MNE; milk N/N intake, g/kg) estimated by mixed-model regression analysis ($MNE = A + BX_1 + CX_2 + DX_3$) of the flow data set derived from 20 metabolic studies in lactating cows where the concentration of both MUN and rumen ammonia nitrogen were determined ($n = 309$)¹

X_1	X_2	X_3	A^2	SE	B^2	SE	C	SE	P -value	D^2	SE	Residual	AICc ³
MY			178	8.7	4.3	0.32						491	2,982
RAN			331	7.7	-4.1	0.73						723	3,094
MUN			327	10.2	-3.3	0.89						762	3,110
RAN	MUN		342	10.3	-3.6	0.78	-1.5	0.94	0.10			721	3,094
MY	RAN		215	9.7	4.2	0.31	-3.6	0.53	<0.001			453	2,941
MY	MUN		219	9.5	4.7	0.30	-5.1	0.65	<0.001			482	2,932
MY	RAN	MUN	233	9.9	4.5	0.30	-2.4	0.56	<0.001	-3.8	0.70	451	2,916

¹MY = milk yield, kg/d; RAN = rumen ammonia N concentration, mg/dL; MUN is in mg/dL.²All P -values ≤ 0.001 .³AICc = Akaike's information criterion with correction.

Table 9. Influence of animal factors on urinary N (UN) excretion (g/d) in lactating cows estimated by mixed-model regression analysis ($UN = A + BX_1 + CX_2 + DX_3$) of the production data set derived from 10 milk-production trials ($n = 443$)¹

X_1	X_2	X_3	A	SE	P -value	B^2	SE	C^2	SE	D^2	SE	Residual	AICc ³
MY			108	11.0	<0.001	2.79	0.336					492	4,301
MUN			126	7.6	<0.001	5.57	0.574					567	4,301
MY	MUN		49	11.1	<0.001	2.68	0.309	5.27	0.534			490	4,234
MY	BW		32	12.3	0.01	1.66	0.317	0.186	0.0173			362	4,204
MY	MUN	DIM	-4	14.8	0.77	3.44	0.339	5.78	0.520	0.205	0.0423	487	4,214
MY	MUN	BW	-27	12.5	0.04	1.59	0.293	4.87	0.473	0.194	0.0167	342	4,124
DMI	MUN	BW	-29	12.3	0.03	4.27	0.669	4.32	0.462	0.136	0.0200	314	4,112

¹MY = milk yield, kg/d; MUN is in mg/dL; BW is in kg/d; DMI is in kg/d.

²All P -values ≤ 0.001 .

³AICc = Akaike's information criterion with correction.

(Table 9). One possible explanation is that increased milk protein concentration lowers the amount of glucose required for the synthesis of other milk components.

In the present evaluation, the influence of DIM on MUN concentration was quadratic with a peak at 120 and 132 d for primiparous and multiparous cows, respectively. A similar quadratic association with a peak between the second and third month of lactation was reported for commercial herds (Stoop et al., 2007). In contrast, the influence of stage of lactation on MUN concentration has been reported to mirror that for milk yield in primiparous cows (Bastin et al., 2009).

Efficiency of Nitrogen Utilization and Milk Urea Concentration

Use of individual cow data indicated that increases in MUN were associated with a decrease in MNE, a

finding consistent with that based on treatment means (Nousiainen et al., 2004). However, a mean decrease of 1.57 g/kg in MNE per mg/dL increase in MUN concentration is considerably lower than a corresponding value of 7.3 g/kg reported previously (Nousiainen et al., 2004). Evaluation of treatment means has shown that dietary CP concentration is the main determinant of MNE in lactating cows (Huhtanen and Hristov, 2009). The current analysis examined the influence of between-cow variation on the relationship between MNE and MUN concentration based on measurements for individual cows using a mixed model that allowed the effect of diet and period to be removed, i.e., for cows fed the same diet at the same time. Higher dietary N intakes are known to increase exponentially the proportion of N excreted in urine (Kebreab et al., 2002). At a constant DMI, more than 80% of the incremental increase in N intake was lost in urine (Huhtanen et al.,

Table 10. Influence of animal factors on the association between rumen ammonia N (RAN) concentration and total-tract OM digestibility (OMD; $n = 414$) and NDF digestibility (NDFD; $n = 414$) or on the association of MUN concentration with OMD ($n = 301$) estimated by mixed-model regression analysis (OMD and $NDFD = A + BX_1 + CX_2 + DX_3$) of the flow data set derived from 26 metabolic studies in lactating cows¹

X_1	X_2	X_3	A^2	SE	B^3	SE	C^2	SE	D^2	SE	Residual	AICc ⁴
OMD												
RAN			725	4.5	1.4	0.42					347	3,872
RAN	RAN \times RAN		711	6.3	4.4	1.49	-0.14	0.042			351	3,867
RAN	DMI		785	10.5	1.4	0.41	-3.1	0.49			307	3,835
RAN	RAN \times RAN	DMI	773	11.0	4.7	1.01	-0.16	0.041	-3.2	0.49	308	3,827
NDFD												
RAN			610	8.0	3.9	0.65					750	4,258
RAN	RAN \times RAN		590	10.7	8.2	1.6	-0.20	0.065			729	4,253
RAN	DMI		677	16.9	4.0	0.62	-3.5	0.78			737	4,241
RAN	RAN \times RAN	DMI	657	17.7	8.8	1.64	-0.22	0.065	-3.7	-0.78	709	4,233
OMD												
MUN			718	6.6	1.8	0.73					364	2,823
MUN	MUN \times MUN		684	13.2	8.9	2.48	-0.34	0.112			373	2,817
MUN	DMI		770	13.9	1.8	0.71	-2.7	0.63			336	2,807
MUN	MUN \times MUN	DMI	737	18.2	8.1	2.45	-0.30	0.111	-2.6	0.62	342	2,803

¹RAN concentration = mg/dL; DMI is in kg/d; MUN is in mg/dL.

²All P -values ≤ 0.001 .

³All P -values ≤ 0.02 .

⁴AICc = Akaike's information criterion with correction.

2008), which explains the strong negative relationship between MNE and MUN concentration. Systematic addition of animal factors (milk yield, BW, and DIM) decreased the residual variance in predictions of MNE from MUN concentration. This suggests that the relationship between MNE and MUN concentration was influenced by factors related to the contribution for maintenance versus productive requirements or nutrient repartitioning. Even though a strong relationship existed between MUN concentration and MNE, it remains unclear whether measurements of MUN concentration can be used to reliably rank individual cows for MNE under commercial conditions. The estimate of the Cow(Exp) variance component for MUN concentration corresponded to a standard deviation value of 1.58 mg/dL (Table 3), whereas the coefficients of MUN concentration averaged -2.53 g of milk N/kg of N intake in the multivariate models (Table 7). Thus, for each 1.58-mg/dL increase in MUN concentration, MNE had a concomitant decrease of 4.0 g/kg, a value equivalent to only 1.4% of the mean MNE determined in the milk-production trials (Table 1). Whereas a moderate heritability for MUN concentration of between 0.13 and 0.22 has been reported (Mitchell et al., 2005; Stoop et al., 2007; König et al., 2008; Bastin et al., 2009), the breeding value for MUN does not appear to be related to the efficiency of N utilization (Vallimont et al., 2011). Selecting for MUN concentration does not appear a useful phenotyping tool for improving MNE, but measurements of MUN concentration on a herd basis allow fine tuning of the diet for improving digestibility or MNE. It is difficult to adjudge whether measurements of MUN concentration for individual cows can be used to reliably rank cows for efficient N utilization.

In the flow data set, the statistical model including both MUN and RAN concentrations explained more of the variation in MNE compared with either variable alone based on AICc calculations. Concentrations of RAN and MUN and excretion of UN are decreased in cows fed diets of lower ruminal protein degradability. Furthermore, increases in absorbed AA in excess of requirements for milk protein or body-tissue synthesis are catabolized and converted to urea in the liver. This may well explain the close association between the concentrations of MUN and dietary CP across a wide range of diets, despite differences in ruminal or host-animal N metabolism. For example, replacing grass silage with red-clover silage resulted in a more efficient utilization of N in the rumen and smaller changes in RAN concentration than would be expected based on the higher dietary CP concentration, but the increase in NAN flow at the omasum was not used for milk protein synthesis (Vanhatalo et al., 2009).

UN Excretion

In the absence of direct measurements of urinary output for the 10 milk-production trials where total-tract digestibility was determined, UN excretion was calculated based on estimates of N retention derived from calculated ME balance (Table 5). Although the lack of direct observations may be criticized, ample evidence exists that N balance studies may overestimate N retention in lactating dairy cows. For example, a meta-analysis based on data from 35 studies reported a mean N balance of $+39$ g/d (Spanghero and Kowalski, 1997). A more recent appraisal across 123 diets indicated a mean N balance of $+30$ g/d (Spek et al., 2013). Assuming a mean muscle CP concentration of 160 g/kg, a net daily N accretion of 30 g/d corresponds to a 1.17 kg/d gain of lean body mass. Several factors may contribute to errors in the determination of N balance including ammonia N losses from feces and urine, excretion of N in urine as nitrate that is not recovered in Kjeldahl analysis, and N losses in scurf (Spanghero and Kowalski, 1997; Reynolds and Kristensen, 2008). It is possible that N balance calculated from estimated ME balance may result in more accurate estimates of UN excretion. However, the approach used in the present evaluation does not take into account variation in feed efficiency that can introduce errors in the estimation of ME balance. Both the residual variance (342 vs. 404) and AICc (4,124 vs. 4,174) for UN excretion decreased using estimated N retention rather than assuming a zero N balance in a 3-variable (milk yield, MUN concentration, and BW; Table 9) model, which would tend to suggest that estimated N balance was correlated with true N retention. For all other models tested (Table 9), a better fit of the data was obtained when N retention was estimated from calculated ME balance compared with estimates based on the assumption of zero N balance.

For individual cows, UN excretion increased 5.8 g/d per 1 mg/dL increase in MUN concentration (Table 9), an estimate considerably lower than values between 12.8 to 14.1 g/d per 1 mg/dL change in MUN concentration estimated using models developed from treatment means (Nousiainen et al., 2004; Spek et al., 2013). Linear relationships between MUN concentration and UN output were found to differ between lactating Holstein and Jersey cows (Kauffman and St-Pierre, 2001). The influence of breed was attributed to differences in BW, and they developed a simple model using $\text{MUN} \times \text{BW}$ to predict UN output. In the present investigation, multivariate models with milk yield and BW as independent factors were used in addition to MUN concentration to estimate UN excretion. Both BW and

milk yield are biologically relevant. Increased milk yield requires higher intakes and greater metabolism of N in body tissues. Maintenance protein requirements that are excreted as N in urine increase with BW (NRC, 2001). Even though between-cow differences were significant for UN output (Table 3), between-cow variability as adjudged from the estimate of the Cow(Exp) variance component corresponded to only 8.3% of the total variance (Table 3). Estimates of the Cow(Exp) variance component for MUN concentration corresponded to a standard deviation value of 1.58 mg/dL (Table 3), whereas the coefficients for MUN averaged 5.1 g of UN per mg/dL-unit change in MUN in the multivariate models (Table 9). Therefore, for each single standard deviation-unit increase in MUN, UN output had a concomitant increase of 8.0 g/d equivalent to 4.1% of mean UN output (Table 5). Compared with the contributions of Exp (50.7%) and Diet(Exp) (30.7%) to total variance in UN output, the extent of variation due to between-animal differences of 8.3% is relatively minor. Across a wide range of diets, MUN concentrations are positively associated with UN output (Jonker et al., 1998; Kauffman and St-Pierre, 2001; Nousiainen et al., 2004). Much less is known about the correlation between the breeding values for MUN concentration with MNE or UN output. The available evidence suggests that the genetic correlation between MUN and MNE is weak (-0.10) and does differ significantly from zero (Vallimont et al., 2011). Given that decreases in MNE are associated with increased UN excretion (Kebreab et al., 2002), it seems reasonable to assume that genetic differences in breeding values for MUN concentration would have little influence on UN output. Given the relatively small extent of variation in UN excretion because of differences between animals, the selection of cows with an inherently low MUN concentration might not be an efficient means for lowering N losses into the environment. Optimizing feeding and management practices appears to offer much more potential for lowering the environmental impact of milk production.

Microbial Nitrogen Supply and Diet Digestibility

This is the first attempt to evaluate the association of between-animal differences in RAN and MUN concentrations with nutrient digestion in lactating cows. Within the range of values in the data sets examined, no association of between-animal differences in RAN concentration with microbial protein synthesis was found, whereas a positive relationship was identified with variability in RAN concentration and total-tract OMD (Table 10). Comparison of these associations suggests that the RAN concentration for optimal digestion is higher than that required for optimal microbial

synthesis. Analysis of treatment means (Oldham, 1984; Nousiainen et al., 2009) has shown that increases in dietary CP concentration by replacing energy supplements with protein supplements were associated with improvements in diet digestibility. However, mechanisms other than increases in RAN concentration might explain the improvement in diet digestion to increases in dietary CP concentration, including the stimulation of cellulolytic bacterial activity by preformed AA and peptides (Atasoglu et al., 2001). Results of in vivo, in situ, and in vitro studies of urea supplementation have been unable to define the RAN requirements of rumen microorganisms to maximize ruminal digestion. Diet digestibility was improved through increases in RAN concentration by urea supplementation in vivo (Kang-Meznarich and Broderick, 1981; Balcells et al., 1993), in situ (Mehrez et al., 1977; Erdman et al., 1986), and in vitro (Wallace, 1979). Evaluation of the flow data set indicated that the positive association between RAN concentration and total-tract OMD and NDFD was not related to intake level, because the regression coefficient for RAN concentration did not change when DMI was included in the model. The negative influence of DMI on total-tract OMD was similar to that reported based on the analysis of treatment means (Huhtanen et al., 2009). Earlier reports concluded that RAN concentrations required to maximize digestion are at least as high as those required for maximum microbial synthesis, and that the optimal concentrations depend on the fermentability of the diet (Schwab et al., 2005). On the basis of published data and the findings from this evaluation, it appears that increasing RAN concentrations have a positive effect on digestibility, irrespective of whether the changes are due to urea supplementation or arise from differences between animals. In the flow data set, the improvement in total-tract OMD with increased RAN concentration was explained entirely by increases in NDFD ($3.9 \times 0.399 = 1.56 \sim 1.40$).

The reasons for the relatively large between-animal variation in RAN concentration ($CV = 0.14$; Table 4) are not clear. Variance may be related to differences in rumen microbial populations, the relative size of protozoal communities in particular. A quantitative meta-analysis of data from studies involving defaunation indicated that concentrations of RAN were markedly higher in faunated than defaunated animals (168 vs. 117 mg of $\text{NH}_3\text{-N/L}$; Eugène et al., 2004). Furthermore, the difference in RAN concentration due to defaunation was also associated with a higher total-tract OM (675 vs. 659 g/kg) and NDF (611 vs. 554 g/kg) digestibility (Eugène et al., 2004). Large between-animal variation in rumen protozoal populations have been reported for sheep fed the same diet (Clarke et al., 1982), but it is not known whether between-animal variation in

protozoal populations has the same influence on RAN concentration and digestibility compared with complete defaunation. Nevertheless, RAN concentration had a negative ($P < 0.001$) association with the ratio of propionate to butyrate in rumen VFA in the flow data set (data not presented), supporting the hypothesis that between-cow variability in RAN concentration was, at least in part, related to differences in protozoal numbers. Comparisons of faunated compared with defaunated animals also indicated that the presence of protozoa in the rumen was associated with lower molar proportions of propionate (202 vs. 231 mmol/mol) and higher molar butyrate proportions (126 vs. 109 mmol/mol; Eugène et al., 2004).

CONCLUSIONS

Concentrations of MUN and RAN varied considerably between cows when the effects due to diet and period were controlled by a statistical model. Increases in MUN or RAN concentration were negatively associated with the efficiency of N utilization but positively related to UN excretion. Concentration of RAN was positively associated with diet digestibility, but the concentration for optimal diet digestibility appears to be higher than that required for maximizing microbial protein supply. Between-cow differences in RAN concentration, although significant, had no influence on microbial N flow or on the efficiency of microbial protein synthesis. Overall, improvements in management and a closer control over diet composition relative to requirements appear to have greater potential to improve the efficiency of N utilization of lactating cows than selection of cows with an inherently low MUN concentration.

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APPENDIX 1

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APPENDIX 2

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